



IN THE UNITED STATES PATENT AND TRADEMARK OFFICE
BEFORE THE BOARD OF PATENT APPEALS AND INTERFERENCES

In re Patent Application of:

Molian DENG *et al.*

Application Serial No.: 09/920,953

Filed: August 2, 2001

Confirmation No.: 9080

Art Unit: 1634

Examiner: James Martinell

Attorney Docket No.: 16517.286/
38-21(52114)B

Title: Nucleic Acid Sequences from *Chlorella sarokiniana* and Uses Thereof

APPEAL BRIEF UNDER 37 C.F.R. § 41.37

Mail Stop Appeal Brief – Patents
Commissioner for Patents
P.O. Box 1450
Alexandria, VA 22313-1450

Sir:

This is an Appeal from the Final Rejection of claims in the above-captioned patent application. A Notice of Appeal was filed on January 10, 2007. Authorization to charge the official fees for this filing is given in the accompanying transmittal letter.

1. Real Party in Interest

The real party in interest is Monsanto Company, a Delaware corporation with offices at 800 North Lindbergh Boulevard, St. Louis, Missouri 63167.

2. Related Appeals and Interferences

Appellants identified the following judicial proceeding, which may have a bearing on the Board's decision in the present Appeal. On May 27, 2004, the Real Party in Interest in the above-captioned matter filed an appeal to the United States Court of Appeals for the Federal

Circuit (“Federal Circuit”) from a decision by the Board in *In re Fisher*. (U.S. Patent Application Serial No. 09/619,643; BPAI Appeal No. 2002-2046; Federal Circuit Case No. 04-1465). The Federal Circuit’s decision in *In re Fisher* may have a bearing on the Board’s decision with regard to at least one of the grounds of rejection in the present appeal. A copy of the Board’s decision in Appeal No. 2002-2046, *Ex parte Fisher*, 72 U.S.P.Q.2d 1020 (Bd. Pat. App. Int. 2004), and a copy of *In re Fisher*, 421 F.3d 1365 (Fed. Cir. 2005) are attached hereto in the Related Proceedings Appendix.

Appellants filed an Appeal Brief in U.S. Patent Application Serial No. 09/684,016, in U.S. Patent Application Serial No. 10/361,942; and in U.S. Patent Application Serial No. 09/199,129, which also may have a bearing on the present appeal. Copies of the Appeal Briefs filed in these applications are also attached hereto in the Related Proceedings Appendix.

3. Status of Claims

Claims 1 to 4 and 6 to 15 are pending. Claim 5 was withdrawn from consideration by the Examiner pursuant to 37 C.F.R. § 1.142(b) in the Office Action mailed March 10, 2004 and was canceled by Appellants in the response dated June 9, 2004. Claims 1 to 4 and 6 to 15 stand finally rejected under 35 U.S.C. § 101 and under 35 U.S.C. § 112, first paragraph. Appellants appeal the rejections of claims 1 to 4 and 6 to 15.

4. Status of Amendments

Appellants have not filed any responses to the Final Office Action dated October 10, 2006 (“Final Action”).

5. Summary of Claimed Subject Matter

Independent Claim 1: The subject matter of independent claim 1 is directed to an isolated nucleic acid molecule comprising a nucleotide sequence of SEQ ID NO: 2 or complement thereof. *Specification* at page 3, lines 3 to 5.

Independent Claim 6: The subject matter of independent claim 6 is directed to a transformed cell having an exogenous nucleic acid molecule which comprises: (A) an exogenous promoter region which functions in said cell to cause the production of a mRNA molecule; which is operably linked to (B) a structural nucleic acid molecule, wherein said structural nucleic acid molecule comprises a nucleic acid sequence of SEQ ID NO: 2 or complement thereof; which is operably linked to (C) a 3' non-translated sequence that functions in said cell to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of said mRNA molecule. *Specification* at page 3, lines 16 to 22.

Independent Claim 11: The subject matter of independent claim 11 is directed to a substantially purified nucleic acid molecule consisting of a nucleic acid sequence of SEQ ID NO: 2 or complement thereof. *Specification* at page 6, lines 34 to 35.

Independent Claim 12: The subject matter of independent claim 12 is directed to a substantially purified nucleic acid molecule comprising a nucleic acid sequence having between 100% and 90% sequence identity with a nucleic acid sequence of SEQ ID NO: 2 or complement thereof. *Specification* at page 7, lines 1 to 3.

A copy of the claims on appeal is attached hereto in the Claims Appendix.

6. Grounds of Rejection to be Reviewed on Appeal

The grounds of rejection to be reviewed in this Appeal are that pending claims 1 to 4 and 6 to 15 stand rejected under 35 U.S.C. § 101 because the claimed invention allegedly lacks patentable utility, and that claims 1 to 4 and 6 to 15 stand rejected under 35 U.S.C. § 112, first paragraph as allegedly failing to comply with the enablement requirement.

7. Argument

A. Summary of Appellants' Position

The specification provides a specific, substantial, and well-established utility for, at a minimum, the nucleic acid sequence of SEQ ID NO: 2, which is an element of all the pending claims, and transformed cells comprising SEQ ID NO: 2. Appellants have provided a statistically significant correlation between the nucleic acid sequence of SEQ ID NO: 2 and a known protein. The utility of the known protein is well-established and the correlation between the known protein and SEQ ID NO: 2 is specific. In setting forth a reasonable correlation between the known protein and the claimed nucleic acid sequence of SEQ ID NO: 2, Appellants have demonstrated that the claimed invention has patentable utility. In other words, Appellants have satisfied the requirements of 35 U.S.C. §§ 101 and 112, first paragraph.

B. The Claimed Nucleic Acids Have Utility under 35 U.S.C. § 101

The Examiner rejected claims 1 to 4 and 6 to 15 under 35 U.S.C. § 101, because the claimed invention allegedly “lacks patentable utility”. Final Action at page 2.

The specification clearly discloses that SEQ ID NO: 2 can be used to encode a 60S Ribosomal Protein L10 or fragment thereof. *Specification* at page 9, lines 15 to 17 and page 68 (Table 1). Upon reading the present specification, one of ordinary skill in the art would readily

recognize that the claimed nucleic acid molecules and cells have utility, for example, to encode a 60S Ribosomal Protein L10 or to modify the expression of 60S Ribosomal Protein L10 in a transformed cell. These utilities are immediately apparent for the claimed nucleic acid molecules, and transformed cells comprising the claimed nucleic acid molecules without further research.

The Examiner has not disputed the use of SEQ ID NO: 2 in encoding a 60S Ribosomal Protein L10. *See*, Office Action mailed March 16, 2006 at page 4 (“Non-final Action”); Final Action at page 2. Rather, the Examiner states that “applicants confuse the identification of a function of a protein with a patentable utility under 35 U.S.C. § 101.” Non-final Action at page 4. Further, the Examiner states that “[i]t is not seen how or where *In re Fischer* [sic] ... creates a *per se* rule that establishes knowledge of a given sequence encoding a known polypeptide to be sufficient to met [sic] the utility requirement under 35 U.S.C. § 101.” Final Action at page 2 (italics in original). This statement misses the point.

In *In re Fisher*, the Federal Circuit reiterated that the “basic *quid pro quo* contemplated by the Constitution and the Congress for granting a patent monopoly is the benefit derived from the public from an invention with *substantial utility*.” *In re Fisher*, 421 F.3d 1365, 1371 (Fed. Cir. 2005) (citing *Brenner v. Manson*, 383 U.S. at 534-35, 1966) (emphasis in original). The Court noted that since “*Brenner* our predecessor court, the Court of Customs and Patent Appeals, and this court have required a claimed invention to have a specific and substantial utility to satisfy § 101.” *Id.* Furthermore, an invention need only provide one identifiable benefit to satisfy 35 U.S.C. § 101. *See Raytheon Co. v. Roper Corp.*, 724 F.2d 951, 958 (Fed. Cir. 1983)

(“when a properly claimed invention meets at least one stated objective, utility under section 101 is clearly shown”).

Although the Supreme Court has not defined the meaning of the terms “specific” and “substantial”, the Federal Circuit has identified a framework for the kind of disclosure an application could contain to establish a specific and substantial utility. *In re Fisher*, 421 F.3d at 1371. First, the Court indicated that to provide a substantial utility, the specification should disclose a utility such that “one skilled in the art can use a claimed discovery in a manner which provides some *immediate benefit to the public*.” *Id.* (emphasis in original). Second, a specific utility can be disclosed by discussing “a use which is not so vague as to be meaningless,” that is that the claimed invention “can be used to provide a well-defined and particular benefit to the public.” *Id.*

Appellants respectfully submit that the question is not whether *In re Fisher* does or does not create a *per se* rule that establishes knowledge of a given sequence encoding a known polypeptide to be sufficient to meet the utility requirement under 35 U.S.C. § 101. Rather the question is whether, in accordance with the ruling in *In re Fisher*, the utility is specific, substantial, and well-established. By showing that SEQ ID NO: 2 can be used to encode a 60S Ribosomal Protein L10 or fragment thereof, a fact that is not contested by the Examiner, Appellants have established that the utility of SEQ ID NO: 2, and transformed cells comprising SEQ ID NO: 2, is specific, substantial, and well-established.

First, the utility of SEQ ID NO: 2 is *specific* because it is specific to SEQ ID NO: 2 and not generally to any nucleic acid sequence. In other words, 60S Ribosomal Protein L10 is not encoded by any general nucleic acid sequence. Second, the utility of SEQ ID NO: 2 is

substantial because the specification discloses that the nucleic acid molecules of the present invention can be used to encode 60S Ribosomal Protein L10 or fragments thereof. *Specification* at page 9, lines 15 to 17, at page 68 (Table 1), and in the sequence listing. In addition, the specification provides for the use of the nucleic acid molecules of the present invention in identifying polymorphisms related to 60S Ribosomal Protein L10. *Id.* at page 53, line 27 through page 59, line 33.

Yet another utility is the use of the nucleic acid molecules of the present invention in transforming plants to modify the expression of 60S Ribosomal Protein L10. *Specification* at page 18, line 25 through page 30, line 12. The specification also discloses that the claimed nucleic acid molecules can also be used in determining the level or pattern of expression of the 60S Ribosomal Protein L10 or mRNA associated with that 60S Ribosomal Protein L10 nucleic acid molecule, for example in a cell. *Id.* at page 52, line 24 through page 53, line 10. These utilities are substantial.

Finally, the utility of SEQ ID NO: 2 is *well-established* because 60S Ribosomal Protein L10 is a well-known, established protein. *See, e.g.,* http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?db=gene&cmd=Retrieve&dopt=full_report&list_uids=6134 (accessed on February 22, 2007) (providing a brief description of this protein).

The Office must accept a stated utility by an applicant unless the Office has evidence or sound scientific reasoning to rebut the applicant's assertion. *In re Oetiker*, 977 F.2d 1443, 1445 (Fed. Cir. 1992). "More specifically, when a patent application claiming a nucleic acid asserts a specific, substantial, and credible utility, and bases the assertion upon homology to existing nucleic acids or proteins having an accepted utility, the asserted utility must be accepted by the

examiner unless the Office has sufficient evidence or sound scientific reasoning to rebut such an assertion.” Federal Register 66(4):1096, Utility Guidelines (2001). Furthermore, “a ‘rigorous correlation’ need not be shown in order to establish practical utility; ‘reasonable correlation’ is sufficient.” *Fujikawa v. Wattanasin*, 93 F.3d 1559, 1565 (Fed. Cir. 1996).

The Office “has the initial burden of challenging a presumptively correct assertion of utility in the disclosure.” *In re Brana*, 51 F.3d 1560, 1567 (Fed. Cir. 1995). The utilities asserted in the specification must be accepted as factually sound unless the Office cites information that undermines the credibility of the assertion. *Id.* The Examiner “must do more than merely question operability – [he] must set forth factual reasons which would lead one skilled in the art to question the objective truth of the statement of operability.” *In re Gaubert*, 524 F.2d 1222, 1224-25 (C.C.P.A. 1975) (emphasis in original); M.P.E.P. § 706.03(a)(1) (“Office personnel are reminded that they must treat as true a statement of fact made by an applicant in relation to an asserted utility, unless countervailing evidence can be provided...”). Here, the Examiner has not met this burden.

The Examiner has provided no support for the assertion that the utilities of SEQ ID NO: 2 are not specific, substantial, and credible. Rather, the entirety of the Examiner’s position is that “(a)pplicants’ assertion that merely knowing that a given sequence encodes a known polypeptide is sufficient to satisfy 35 U.S.C. § 101 is not convincing” because, according to the Examiner, *In re Fisher* suggests that mere knowledge that a given sequence encodes a known polypeptide is insufficient to satisfy the utility requirement under 35 U.S.C. § 101. Final Action at page 2.

The Examiner has set forth neither scientific publications nor factual reasons that undermine the credibility of Appellants’ assertion that the utilities of SEQ ID NO: 2 are specific,

substantial, and credible. In fact, as noted previously, the Examiner admits that SEQ ID NO: 2 can be used to encode a 60S Ribosomal Protein L10 or fragment thereof. As Appellants have pointed out, the fact that SEQ ID NO: 2 can be used to encode a 60S Ribosomal Protein L10 or fragment thereof is evidence that the utility is specific, substantial, and well-established. Therefore, the claimed invention meets the utility test set forth in *In re Fisher*.

In conclusion, Appellants respectfully submit that the claimed invention has specific, substantial, and well-established utility and request that the Board reverse the rejection of claims 1 to 4 and 6 to 15 under 35 U.S.C. § 101.

C. The Claimed Nucleic Acids Satisfy 35 U.S.C. § 112

The Examiner rejected claims 1 to 4 and 6 to 15 under 35 U.S.C. § 112, first paragraph, because the claimed invention allegedly fails to comply with the enablement requirement. The Examiner states that “[s]ince the claimed invention lacks utility under 35 U.S.C. § 101, the instant application does not teach how to use the invention”. Final Action at page 2.

Appellants submit that this rejection has been overcome by the arguments set forth above with respect to the rejection under 35 U.S.C. § 101. In other words, Appellants respectfully submit that the claimed invention has specific, substantial, and well-established utility and request that the Board reverse the rejection of claims 1 to 4 and 6 to 15 under 35 U.S.C. § 112, first paragraph.

CONCLUSION

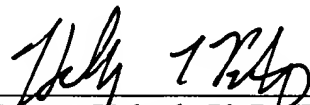
In view of the foregoing, Appellants respectfully request that the Board of Patent Appeals and Interferences reverse the pending rejections and that the subject application be allowed forthwith.

Respectfully submitted,

Date: March 12, 2007

CORRESPONDENCE ADDRESS:

Monsanto Company
800 N. Lindbergh Blvd.
Attn: Gail P. Wuellner, IP Paralegal
IP Paralegal (E2NA)
St. Louis, Missouri 63167



Gautam Prakash, Ph.D. (Reg. Agent No. 53,481)
Holly L. Prutz (Reg. Attorney No. 47,755)
David R. Marsh (Reg. Attorney No. 41,408)
Arnold & Porter LLP
555 Twelfth Street, N.W.
Attn: IP Docketing
Washington, DC 20004
Tel: 202-942-5000
Fax: 202-942-5999

CLAIMS APPENDIX

Claim 1. An isolated nucleic acid molecule comprising a nucleotide sequence of SEQ ID NO: 2 or complement thereof.

Claim 2. The isolated nucleic acid molecule of claim 1, wherein said isolated nucleic acid molecule encodes a *Chlorella sarokiniana* protein.

Claim 3. The isolated nucleic acid molecule of claim 2, wherein said *Chlorella sarokiniana* protein is a homologue of a 60S Ribosomal protein.

Claim 4. The isolated nucleic acid molecule of claim 3, wherein said isolated nucleic acid molecule a nucleotide sequence of SEQ ID: 2.

Claim 6. A transformed cell having an exogenous nucleic acid molecule which comprises:

- (A) an exogenous promoter region which functions in said cell to cause the production of a mRNA molecule; which is operably linked to
- (B) a structural nucleic acid molecule, wherein said structural nucleic acid molecule comprises a nucleic acid sequence of SEQ ID NO: 2 or complement thereof; which is operably linked to
- (C) a 3' non-translated sequence that functions in said cell to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of said mRNA molecule.

Claim 7. The transformed cell according to claim 6, wherein said cell is selected from the group consisting of an algal cell, a plant cell, a mammalian cell, a bacterial cell, a fungal cell and an insect cell.

Claim 8. The transformed cell according to claim 7, wherein said cell is an algal cell.

Claim 9. The transformed cell according to claim 8, wherein said cell is a *Chlorella sarokiniana* cell.

Claim 10. The transformed cell according to claim 7, wherein said cell is a plant cell.

Claim 11. A substantially purified nucleic acid molecule consisting of a nucleic acid sequence of SEQ ID NO: 2 or complement thereof.

Claim 12. A substantially purified nucleic acid molecule comprising a nucleic acid sequence having between 100% and 90% sequence identity with a nucleic acid sequence of SEQ ID NO: 2 or complement thereof.

Claim 13. The substantially purified nucleic acid molecule of claim 12, wherein said nucleic acid molecule comprises a nucleic acid sequence having between 100% and 95% sequence identity with a nucleic acid sequence of SEQ ID NO: 2 or complement thereof.

Claim 14. The substantially purified nucleic acid molecule of claim 13, wherein said nucleic acid molecule comprises a nucleic acid sequence having between 100% and 98% sequence identity with a nucleic acid sequence of SEQ ID NO: 2 or complement thereof.

Claim 15. The substantially purified nucleic acid molecule of claim 14, wherein said nucleic acid molecule comprises a nucleic acid sequence having between 100% and 99% sequence identity with a nucleic acid sequence of SEQ ID NO: 2 or complement thereof.

EVIDENCE APPENDIX

None

RELATED PROCEEDINGS APPENDIX

Copies of the following are attached:

1. *Ex parte Fisher*, 72 U.S.P.Q.2d 1020 (Bd. Pat. App. Int. 2004);
2. *In re Fisher*, 421 F.3d 1365 (Fed. Cir. 2005);
3. Appeal Brief filed in U.S. Patent Application Serial No. 09/684,016;
4. Appeal Brief filed in U.S. Patent Application Serial No. 10/361,942; and
5. Appeal Brief filed in U.S. Patent Application Serial No. 09/199,129.



**IN THE UNITED STATES PATENT AND TRADEMARK OFFICE
BEFORE THE BOARD OF PATENT APPEALS AND INTERFERENCES**

In re application of:

David K. KOVALIC *et al.*

Appln. No.: 09/684,016

Filed: October 10, 2000

Title: **Annotated Plant Genes**

Confirm No: 9497

Art Unit: 1631

Examiner: Shubo ZHOU

Atty. Docket: 16517.031

APPELLANTS' FOURTH AMENDED BRIEF

Mail Stop Appeal Brief – Patents

Commissioner for Patents

P.O. Box 1450

Alexandria, Virginia 22313-1450

Sir:

This is an Appeal from the Final Rejection of all claims pending in the above-referenced patent application. A Notice of Appeal was filed on September 17, 2003. An Appellant's Brief was filed November 17, 2003, at which time the statutory fee of \$320.00 for submitting an appeal brief was paid. A Second Amended Brief is submitted in response to the Office Communication mailed April 7, 2005 which alleged that the Brief filed August 28, 2003, and Amended Brief filed April 1, 2004 was non-compliant with 37 C.F.R. 1.192(c). A Third Amended Brief was submitted in response to the Office Communication mailed June 1, 2005 which alleged that the Brief filed May 6, 2005, was non-compliant with 37 C.F.R. 1.192(c). This Fourth Amended Brief is submitted in response to the Office Communication mailed November 24, 2006, which alleged that the Brief filed July 1, 2006 was non-compliant with 37 C.F.R. 41.37(c). This Fourth Amended Brief is filed pursuant to the format set forth in 37 C.F.R. § 41.37.

1. Real Party in Interest

The real party in interest is Monsanto Company, a Delaware corporation with offices at 800 North Lindbergh Boulevard, St. Louis, Missouri 63167.

2. Related Appeals and Interferences

Appellants identify the following judicial proceeding, which may have a bearing on the Board's decision in the present Appeal. On May 27, 2004, the Real Party in Interest in the above-captioned matter filed an appeal to the United States Court of Appeals for the Federal Circuit ("Federal Circuit") from a decision by the Board in *In re Fisher*. (U.S. Appln. No. 09/619,643, B.P.A.I. Appeal No. 2002-2046, Fed. Cir. Case No. 04-1465). The Federal Circuit's decision in *In re Fisher* may have a bearing on the Board's decision with regard to at least one of the grounds of rejection in the present appeal. A copy of the Board's decision in Appeal No. 2002-2046 is attached hereto in the "Related Proceedings Appendix."

In addition, Appellants also identify the following additional Board decisions which may have a bearing on the instant appeal: U.S. Appln. No. 09/654,617, B.P.A.I. Appeal No. 2003-1744; U.S. Appln. No. 09/620,392, B.P.A.I. Appeal No. 2003-1746; U.S. Appln. No. 09/540,232, B.P.A.I. Appeal No. 2003-1137; U.S. Appln. No. 09/440,687, B.P.A.I. Appeal No. 2003-1504; U.S. Appln. No. 09/565,240, B.P.A.I. Appeal No. 2003-1135; U.S. Appln. No. 09/540,215, B.P.A.I. Appeal No. 2003-0996; U.S. Appln. No. 09/552,087, B.P.A.I. Appeal No. 2004-1772; and U.S. Appln. No. 09/206,040, B.P.A.I. Appeal No. 2002-0078. Copies of the Board's decisions in these Appeals are also attached hereto in Appendix B.

3. Status of Claims

Claims 11-16 are pending. Claims 1-10 were cancelled in a Response filed March 11, 2002. Claims 11-16 stand finally rejected under 35 U.S.C. § 101 as allegedly lacking utility and under 35 U.S.C. § 112, first paragraph, as allegedly lacking enablement. Claims 11-15 are rejected under 35 U.S.C. § 112, first paragraph, as allegedly lacking written description. Claim 14 stands rejected under 35 U.S.C. § 112, first paragraph, as allegedly containing new matter. Claim 13 is rejected under 35 U.S.C. § 102(b), as allegedly being anticipated. Appellants appeal all of the rejections of claims 11-16. A copy of the claims on appeal is provided in the "Claims Appendix" attached hereto.

4. Status of Amendments

Appellants filed an Amendment After Final Rejection ("Amendment") on July 29, 2003, requesting amendment of claims 13-15. The Amendment was filed in response to the Final Office Action ("Final Action"), which was mailed on June 17, 2003 (Paper No. 15). In response to Appellants' Amendment, an Advisory Action was mailed by the U.S. Patent and Trademark Office on August 27, 2003 (Paper No. 18) ("Advisory Action"), stating that "[f]or purposes of Appeal, the proposed amendment(s) will be entered...."

5. Summary of the Claimed Subject Matter

The claimed subject matter of independent claim 11 is directed to a substantially purified nucleic acid molecule comprising a fragment nucleic acid molecule having from about 30 to about 50 nucleotide residues of a nucleic acid molecule having the nucleotide sequence of SEQ ID NO: 48411. Specification at page 3, lines 12-14; page 8, line 28 through page 9, line 9.

The claimed subject matter of independent claim 12 is directed to a substantially purified nucleic acid molecule comprising a fragment nucleic acid molecule having from about 50 to about 100 nucleotide residues of a nucleic acid molecule having the nucleotide sequence of SEQ ID NO: 48411. *Id.*

The claimed subject matter of independent claim 13 is directed to a substantially purified nucleic acid molecule comprising a fragment nucleic acid molecule having from about 30 to about 50 nucleotide residues, wherein said fragment nucleic acid molecule exhibits complete complementarity to a fragment of a second nucleic acid molecule having the nucleotide sequence of SEQ ID NO: 48411 or a complete complement thereof. Specification at page 3, lines 12-18; page 8, line 28 through page 9, line 9; page 9, line 30 through page 10, line 4.

The claimed subject matter of independent claim 14 is directed to a substantially purified nucleic acid molecule having between 90% and 100% sequence identity with nucleotides 1 through 123 of SEQ ID NO: 48411 or a complete complement thereof.

Specification at page 3, lines 12-18; page 11, line 5 through page 12, line 12; and sequence listing at SEQ ID NO: 48411.

6. Grounds of Rejection to be Reviewed on Appeal

The grounds of rejection to be reviewed in this Appeal are:

- (a) pending claims 11-16 stand rejected under 35 U.S.C. § 101, for allegedly not being supported by a specific asserted utility or a well established utility;
- (b) pending claims 11-16 stand rejected under 35 U.S.C. § 112, first paragraph for alleged lack of enablement because the claimed invention purportedly lacks utility;
- (c) pending claims 11-15 stand rejected under 35 U.S.C. § 112, first paragraph for alleged insufficiency of written description;
- (d) pending claim 14 stands rejected under 35 U.S.C. § 112, first paragraph for allegedly containing new matter; and
- (e) pending claim 13 stands rejected under 35 U.S.C. § 102(b) for alleged anticipation.

A. Grouping of Claims

Claims 11-16 are pending in this application. All of the claims at issue do not stand or fall together. The separate patentability of claims 11-16 is addressed together in Sections 8.A through 8.C below. The separate patentability of claims 11-15 is addressed in Section 8.D below. The separate patentability of claim 14 is addressed in Section 8.E below. The separate patentability of claim 13 is addressed in Section 8.F below.

7. Preliminary Remarks

Appellants thank the Examiner for withdrawing the rejection of claim 15 under 35 U.S.C. §112, second paragraph in the Advisory Action at page 2.

8. Argument

A. Summary of Appellants' Position

As the Supreme Court said in *Brenner v. Manson*, the "basic quid pro quo contemplated by the Constitution and the Congress for granting a patent monopoly is the

benefit derived by the public from an invention with substantial utility....where specific benefit exists in currently available form.” 383 U.S. 519, 534-35, 148 U.S.P.Q. 689, 695 (1966). Appellants have met their part of the bargain – they have disclosed nucleic acid molecules that, in their current form, provide at least one specific benefit to the public, for example, use to identify the presence or absence of a polymorphism. This benefit is specific, not vague or unknown, and it is a “real world” or substantial benefit. Because the claimed nucleic acids provide at least this benefit, they satisfy the utility requirement of 35 U.S.C. § 101. Because the specification teaches how to make and use the claimed nucleic acids for the disclosed utilities, the enablement requirement of 35 U.S.C. § 112 has likewise been met.

Furthermore, Appellants have provided an adequate description of the claimed nucleic acids that demonstrates Appellants’ possession of the claimed invention. Each genus of claimed nucleic acid molecules, *e.g.*, the nucleic acid molecules comprising the nucleic acid sequence of SEQ ID NO: 48411 its complement, and fragments thereof, for example, has been described by the recitation of a common structural feature – the nucleotide sequences of SEQ ID NO: 48411, and its complement, respectively – which distinguishes molecules within the claimed genus from molecules outside of the claimed genus. Because the specification demonstrates that Appellants have possession of (and have provided an adequate description of) the claimed genera of nucleic acid molecules, the specification satisfies the written description requirement of 35 U.S.C. § 112.

Appellants have provided sufficient written description support in the specification and in the sequence listing such that a new matter rejection is improperly applied to claim 14. The recitation of the range “nucleotides 1 through 123 of SEQ ID NO: 48411” is supported in the specification and in the sequence listing; and the inclusion of such a range in the presently pending claim is validated by *In re Wertheim*, 541 F.2d 257, 191 U.S.P.Q. 90 (C.C.P.A. 1976). As such, Appellants have met the burden of written description and introduce no new matter by the inclusion of the noted claim language.

Claim 13 was erroneously rejected as anticipated by a reference that fails to teach the recited nucleic acid sequence. The Examiner improperly considered a non-identical chemical compound to anticipate the claims as drawn to a nucleic acid molecule comprising a fragment nucleic acid molecule having from about 30 to about 50 nucleotide residues, wherein said fragment nucleic acid molecule exhibits complete complementarity to a fragment of a second nucleic acid molecule having the nucleotide sequence of SEQ ID NO: 48411 or a complete complement thereof, despite the fact that the cited reference fails to teach such a fragment nucleic acid molecule. The Examiner has asserted an untenable interpretation of claim 13, misconstruing claim 13 and citing a reference that does not anticipate the present claims. Absent a teaching of each and every element of the claims, the reference cited by the Examiner does not anticipate the present claim 13.

B. The Claimed Nucleic Acids Have Legal Utility

Pending claims 11-16 were erroneously rejected under 35 U.S.C. § 101 because the claimed invention was allegedly not supported by either a “specific and/or substantial utility or a well established utility.” Final Action at pages 2-3. According to the Final Action, “since the function of the gene comprising the claimed sequence is not known, identifying the presence or absence of a polymorphism in a population is not deemed a real world utility.” *Id.*

This analysis misstates the nature of the asserted uses, ignores disclosed utilities, and misapplies the doctrine of “practical utility” developed by the courts after *Brenner v. Manson*. The “threshold for utility is not high: An invention is ‘useful’ under section 101 if it is capable of providing some identifiable benefit.” *Juicy Whip, Inc. v. Orange Bang, Inc.*, 185 F.3d 1364, 1366, 51 U.S.P.Q.2d 1700, 1702 (Fed. Cir. 1999), *citing Brenner v. Manson*, 383 U.S. 519, 534 (1966). Furthermore, an invention need only provide one identifiable benefit to satisfy 35 U.S.C. § 101. *See Raytheon Co. v. Roper Corp.*, 724 F.2d 951, 958, 220 U.S.P.Q. 592, 598 (Fed. Cir. 1983) (“when a properly claimed invention meets at least one stated objective, utility under section 101 is clearly shown”).

The courts have expressed a test for utility that hinges on whether an invention provides an “identifiable benefit.” *Juicy Whip, Inc. v. Orange Bang, Inc.*, 185 F.3d 1364, 1366, 51 U.S.P.Q.2d 1700, 1702 (Fed. Cir. 1999), *citing Brenner v. Manson*, 383 U.S. 519, 534 (1966). For analytical purposes, the requirement for an “identifiable benefit” may be broken into two prongs: (1) the invention must have a specific, *i.e.*, not vague or unknown benefit, *In re Brana*, 51 F.3d 1560, 1565, 34 U.S.P.Q.2d 1436, 1440 (Fed. Cir. 1995); and (2) the invention must provide a real world, *i.e.*, practical or “substantial” benefit. *Fujikawa v. Wattanasin*, 93 F.3d 1559, 1563, 39 U.S.P.Q.2d 1895, 1899 (Fed. Cir. 1996). A corollary to this test for utility is that the invention must not be “totally incapable of achieving a useful result,” *i.e.*, the utility must not be incredible or unbelievable. *Brooktree Corp. v. Advanced Micro Devices, Inc.*, 977 F.2d 1555, 1571, 24 U.S.P.Q.2d 1401, 1412 (Fed. Cir. 1992).

Appellants have asserted throughout the specification that the claimed nucleic acid molecules provide identifiable benefits, for example use to identify the presence or absence of a polymorphism, and use as a marker. *See, e.g.*, specification at page 39, line 29 through page 44, line 2. Either of these utilities alone is enough to satisfy Section 101. Because Appellants need only establish a single utility to satisfy 35 U.S.C. § 101, and they have done so in the present case, the premise of the rejection under Section 101 is incorrect, and the rejection should be reversed.

**(1) The Claimed Nucleic Acid Molecules Provide A Specific Benefit,
i.e., They Have Specific Utility**

Appellants have demonstrated that the claimed nucleic acid molecules are themselves useful for utilities disclosed in the specification, *e.g.*, to detect the presence or absence of polymorphisms. *See, e.g.*, specification at page 40, line 4 through page 42, line 13. The specification also discloses additional utilities for the claimed nucleic acid molecules, including, for example, use of the claimed nucleic acid molecules to measure the

level of mRNA in a sample,¹ and use as molecular markers.² See *e.g.*, specification at page 39, line 29 through page 40, line 3; page 42, line 14 through page 44, line 3; page 44, lines 11-16.

(a) Identifying the Presence or Absence of a Polymorphism

One of the utilities disclosed in the specification is use of the claimed nucleic acid molecules to identify the presence or absence of a polymorphism. Specification at page 40, line 4 through page 42, line 13. The Examiner argues that this utility is not “a real world utility”, *see* Final Action at page 3, but does not provide any support, legal or factual, for the proposition that detection of polymorphisms is not a legal utility. The Examiner’s reliance upon the Interim Utility Guidelines has led to an interpretation of utility that contravenes well-established doctrines of utility developed in the courts.

Appellants reiterate that many of the disclosed utilities in this case, including detection of polymorphisms, are directly analogous to the utilities of a microscope, *i.e.*, the claimed nucleic acid molecules may be used to locate and measure nucleic acid molecules within a sample, cell, or organism. The Examiner denigrates this utility by asserting that these uses are not “useful” because allegedly “...further research has to be done....” See, *e.g.*, Final Action at page 3. However, the fact that, for example, a new and nonobvious microscope or screening assay can be used for further learning about products or processes does not lessen the fact that such “tools” have legal utility. Indeed,

¹ It is standard practice to screen populations of nucleic acids with EST sequences, often attached to a microarray, without characterizing each and every target mRNA. Knowing that the gene corresponding to the claimed nucleic acid molecules is expressed under certain conditions or in certain tissues or at certain levels is in itself useful. For example, such information is useful to detect expression changes in traits of interest.

² One can use the claimed nucleic acid molecules to determine location of a corresponding DNA sequence on a physical map or genetic map location without knowing anything beyond the claimed sequence. The use of molecular markers is a practical activity in the development of nutritionally enhanced or agriculturally enhanced crops. Such markers are useful in, for example, genetic mapping or linkage analysis, marker-assisted breeding, physical genome mapping, transgenic crop production, crop monitoring diagnostics, and gene identification and isolation. As more markers are identified, genetic maps will become more detailed and it will be easier for plant breeders to breed for particular traits.

"Many research tools such as gas chromatographs, screening assays, and nucleotide sequencing techniques have clear, specific and unquestionable utility (*e.g.*, they are useful in analyzing compounds)." MPEP § 2107.01 at page 2100-33.

Use of the claimed nucleic acid molecules to detect the presence or absence of polymorphisms is no more legally insufficient than using a gas chromatograph to analyze the chemical composition of a gas – such use determines information about the gas, not the gas chromatograph. Moreover, even if the gas chromatograph detects the absence of a particular chemical element in the gas, that finding does not obviate the utility of the gas chromatograph itself. Information has been obtained about the gas.³ Likewise, the claimed nucleic acid molecules have utility even if the absence of a particular polymorphism is detected. Indeed, the absence of a polymorphism usefully demonstrates that the two (or more) populations being compared share a common genetic heritage.

The claimed nucleic acid molecules produce a specific, *i.e.*, not vague or unknown, benefit – they are useful to identify the presence or absence of a polymorphism. This benefit is immediately realized directly from the use of the claimed nucleic acids, not from the use of other molecules. Such a proven use, that provides an acknowledged benefit to the public, satisfies the utility requirement of 35 U.S.C. § 101.

(b) Probes for Other Molecules or Source for Primers

Other uses for the claimed nucleic acid molecules include use as probes for other molecules or as a source of primers. The specification discloses that the claimed nucleic acid molecules can be used, via hybridization, in real world applications, such as for example, to isolate nucleic acid homologues of other plants and organisms including alfalfa,

³ For example, gas sampled from crude oil may be analyzed by gas chromatography for the presence or absence of chlorine, which is toxic to catalysts used in gasoline refining even in very low concentrations. The absence of a peak at the molecular weight of chlorine indicates the absence of chlorine in the sample being tested, thereby providing useful information (no chlorine is present, therefore the catalyst will not be destroyed) to the refinery manager. See, *e.g.*, U.S. Patent No. 6,133,740 entitled "Chlorine Specific Gas Chromatographic Detector."

Arabidopsis, barley, *Brassica*, broccoli, cabbage, etc.⁴ Specification at page 38, lines 5-15. The Examiner has not provided any evidence that would reasonably suggest that this cannot be done, and as such, has not met the burden of proof required to establish a utility rejection. See *In re Brana*, 51 F.3d 1560, 1567, 34 U.S.P.Q.2d 1436, 1441 (Fed. Cir. 1995). Accord *In re Gaubert*, 524 F.2d 1222, 1225-26, 187 U.S.P.Q. 664, 666 (C.C.P.A. 1975); *In re Langer*, 503 F.2d 1380, 1391, 183 U.S.P.Q. 288, 297 (C.C.P.A. 1974).

One illustrative example of a molecule that can be isolated using a claimed nucleic acid molecule is the promoter of the gene corresponding to that claimed nucleic acid molecule. Appellants have specifically disclosed that one use of the claimed nucleic acid molecules is to initiate a chromosome walk. See *e.g.*, specification at page 39, lines 4-16. The Examiner denigrates Appellants' disclosed utilities by asserting that they are not "specific." Final Action at page 2-3. In short, the Examiner suggests that the asserted utilities are legally insufficient simply because other molecules can be used for the same purpose, *i.e.*, a chromosome walk. This position is wrong as a matter of law --- there is no requirement of exclusive utility in the patent law. See *Carl Zeiss Stiftung v. Renishaw PLC*, 945 F.2d 1173, 1180, 20 U.S.P.Q.2d 1094, 1100 (Fed. Cir. 1991) ("An invention need not be the best or the only way to accomplish a certain result..."). Such an argument would imply that a new golf club has no legal utility because other golf clubs can be used for the same purpose, *i.e.*, hitting golf balls. That position must be rejected as it requires reading "into the patent laws limitations and conditions which the legislature has not expressed," a practice condemned by the Supreme Court. See *Diamond v. Chakrabarty*, 447 U.S. 303, 308, 206 U.S.P.Q. 193, 196 (1980), quoting *United States v. Dubilier Condenser Corp.*, 289 U.S. 178, 199, 17 U.S.P.Q. 154, 162 (1933).

⁴ Moreover, one skilled in the art of hybridization and amplification understands how to design and utilize probes and primers to target a sequence of interest, and thus it is not necessary for Appellants to provide a laundry list of each and every nucleic acid molecule that can be identified using the claimed nucleic acid molecules.

Moreover, Appellants reiterate that it is factually incorrect that this use is not “specific” to the claimed nucleic acids. The claimed nucleic acid molecules provide a particularly appropriate and demonstrably useful starting point for a walk to isolate a promoter that is active, for example, in *Glycine max*. A random nucleic acid molecule does not provide an equally good starting point to isolate such a promoter. Furthermore, even if another nucleic acid molecule provided a better starting point than the claimed nucleic acid molecules, it would not obviate the utility of the claimed nucleic acid molecules. An invention may be “less effective than existing devices but nevertheless meet the statutory criteria for patentability.” *Custom Accessories, Inc. v. Jeffrey-Allan Indus.*, 807 F.2d 955, 960 n.12, 1 U.S.P.Q.2d 1196, 1199 n.12 (Fed. Cir. 1986).

The Examiner has failed to provide evidence for believing that the claimed nucleic acid molecules could not be so used. Accordingly, the demonstration of utility through use as a probe for other molecules or as a source of primers satisfies the requirements of 35 U.S.C. § 101. *See In re Brana*, 51 F.3d 1560, 1566, 34 U.S.P.Q.2d 1436, 1441 (Fed. Cir. 1995).

(2) The Claimed Nucleic Acid Molecules Provide Practical, Real World Benefits, *i.e.*, They Have Substantial Utility

It appears that the Final Action is arguing that the disclosed uses are legally insufficient or “insubstantial” under 35 U.S.C. § 101, but such an argument has no basis in law. The touchstone of “substantial” utility is “real world” or “practical utility.” *See, e.g., Fujikawa v. Wattanasin*, 93 F.3d 1559, 1563, 39 U.S.P.Q.2d 1895, 1899 (Fed. Cir. 1996). “‘Practical utility’ is a shorthand way of attributing ‘real world’ value to claimed subject matter. In other words, one skilled in the art can use a claimed discovery in a manner which provides some immediate benefit to the public.” *Nelson v. Bowler*, 626 F.2d 853, 856, 857, 206 U.S.P.Q. 881, 883 (C.C.P.A. 1980) (“tests evidencing pharmaco-

logical activity may manifest a practical utility even though they may not establish a specific therapeutic use”).⁵

There can be no question that one skilled in the art can use the claimed nucleic acid molecules in a manner which provides an immediate benefit to the public, for example, to perform high-throughput microarray analysis of expression changes in a series of tissue samples. The detection of expression changes provides an immediate benefit to the public because, for example, it enables a plant geneticist to rapidly identify relationships or patterns within the expression changes corresponding to various tissues of organisms grown under various different conditions. This comparative information about a plant's expression profile under different growth conditions, like the information about a compound's pharmacological profile in *Nelson*, provides an immediate benefit and thus a practical real world utility to the public.

Quite apart from the analysis of gene expression, there is also no question that the public has recognized the benefits provided by the claimed subject matter, and has attributed “real world” value to such nucleic acid molecules. The utility of ESTs is not merely an academic issue; the real world value of ESTs is self-evident from the growth of a multi-million dollar industry in the United States premised on the usefulness of ESTs. Like fermentation processes involving bacteria, ESTs and nucleic acid molecules with EST sequences are “industrial product[s] used in an industrial process – a useful or technical art if there ever was one.” See, e.g., *In re Bergy*, 563 F.2d 1031, 1038, 195 U.S.P.Q. 344, 350 (C.C.P.A. 1977).

The market participants for EST products are primarily sophisticated corporations and highly knowledgeable scientists who are unlikely to pay for useless inventions. Compare *Raytheon Co. v. Roper Corp.*, 724 F.2d 951, 960, 220 U.S.P.Q. 592, 599 (Fed. Cir. 1983) (“People rarely, if ever, appropriate useless inventions”). Quite simply, the

⁵ Accord *Cross v. Iizuka*, 753 F.2d 1040, 1050, 224 U.S.P.Q. 739, 747-48 (Fed. Cir. 1985); *Rey-Bellet v. Engelhardt*, 493 F.2d 1380, 1383, 181 U.S.P.Q. 453, 454 (C.C.P.A. 1974).

commercial value of ESTs is proof of their real world value and of the benefits they provide to the public. This evidence cannot be ignored. The patent system was created to serve and foster growth and development in the industrial arts. If the industries themselves recognize and appreciate the value of an invention, it is not for the Patent Office to say that they are mistaken.

(3) The Disclosed Utilities Are Credible to One of Skill in the Art

An assertion of utility must be accepted by the Examiner unless it would not be considered “credible” by a person of ordinary skill in the art. MPEP § 2107 at 2100-29. Cases in which utility was found not to be credible are rare, and usually involve “hare-brained” utilities.⁶ A challenge to the credibility of a utility is essentially a challenge directed to operability, and such a challenge must be supported by a clear statement of “factual reasons which would lead one skilled in the art to question the objective truth of the statement of operability.” *In re Gaubert*, 524 F.2d 1222, 1225-26, 187 U.S.P.Q. 664, 666 (C.C.P.A. 1975); *see In re Brana*, 51 F.3d 1560, 1567, 34 U.S.P.Q.2d 1436, 1441 (Fed. Cir. 1995); MPEP § 2107.02 at 2100-41.

Appellants have explicitly identified specific and substantial utilities, not only in the specification, but in Applicants’ Response dated August 8, 2002 at page 6, lines 7 through 14. “To violate [35 U.S.C.] 101 the claimed device must be totally incapable of achieving a useful result.” *Brooktree Corp. v. Advanced Micro Devices, Inc.*, 977 F.2d

⁶ Examples of incredible utilities are given in MPEP § 2107.01 at page 2100-34, and include:

an invention asserted to change the taste of food using a magnetic field (*Fregeau v. Mossinghoff*, 776 F.2d 1034, 227 U.S.P.Q. 848 (Fed. Cir. 1985)), a perpetual motion machine (*Newman v. Quigg*, 877 F.2d 1575, 11 U.S.P.Q. 1340 (Fed. Cir. 1989)), a flying machine operating on “flapping or flutter function” (*In re Houghton*, 433 F.2d 820, 167 U.S.P.Q. 687 (C.C.P.A. 1970)), a method for increasing the energy output of fossil fuels upon combustion through exposure to a magnetic field (*In re Ruskin*, 354 F.2d 395, 148 U.S.P.Q. 221 (C.C.P.A. 1966)), uncharacterized compositions for curing a wide array of cancers (*In re Citron*, 325 F.2d 248, 139 U.S.P.Q. 516 (C.C.P.A. 1963)), a method of controlling the aging process (*In re Eltgroth*, 419 F.2d 918, 164 U.S.P.Q. 221 (C.C.P.A. 1970)), and a method of restoring hair growth (*In re Ferens*, 417 F.2d 1072, 163 U.S.P.Q. 609 (C.C.P.A. 1969)).

1555, 1571, 24 U.S.P.Q.2d 1401, 1412 (Fed. Cir. 1992). To date, the Examiner has provided no conclusive evidence that the claimed nucleic acid molecules will not work for the disclosed utilities. Unless and until the Examiner can prove that the claimed invention is wholly inoperative, the rejection must be withdrawn.

In view of the above, Appellants contend that the claimed nucleic acid molecules are supported by credible, specific, and substantial utilities disclosed in the specification. Moreover, the Examiner has failed to raise any credible evidence challenging the presently asserted utilities. Consequently, the rejection of claims 11-16 under 35 U.S.C. §101 is improper and should be reversed.

C. The Claimed Nucleic Acids Are Enabled by the Specification

The enablement of the claimed nucleic acid molecules has been challenged. Claims 11-16 have been erroneously rejected as not enabled by the specification, because the claimed nucleic acid molecules allegedly lack utility and therefore cannot be enabled. Final Action at pages 3-4, Advisory Action at page 2. This rejection is erroneous and has been overcome by the arguments stated above regarding utility because it is well-established law that "the enablement requirement is met if the description enables any mode of making and using the invention." *Johns Hopkins University v. CellPro*, 152 F.3d 1342, 1361, 47 U.S.P.Q.2d 1705, 1719 (Fed. Cir. 1998) (emphasis added), quoting *Engel Indus. v. Lockformer Co.*, 946 F.2d 1528, 1533, 20 U.S.P.Q.2d 1300, 1304 (Fed. Cir. 1991). Unless and until the Examiner comes forth with evidence to rebut the objective truth of the utilities disclosed in the specification, this enablement rejection must be withdrawn as improper. See *In re Wright*, 999 F.2d 1557, 1561-62, 27 U.S.P.Q.2d 1510, 1513 (Fed. Cir. 1993); *Ex parte Lemak*, 210 U.S.P.Q. 306, 307 (Bd. App. 1981) ("pure conjecture" does not substantiate rejection for lack of enablement).

D. The Specification Provides An Adequate Written Description of the Claimed Invention

The adequacy of the written description of the claimed invention has been challenged by the Examiner because the claimed subject matter was allegedly "not described

in the specification in such a way as to reasonably convey to one skilled in the relevant art that the inventor(s)...had possession of the claimed invention.” Final Action at page 4. The Examiner contends that “the specification only provides sequences of the elected SEQ ID NO: 48411, but not the sequences comprising the sequence of the elected SEQ ID NO or comprising a fragment thereof.” Final Action at page 4. This is not a proper basis for a written description rejection of a “comprising” claim. If it were, every “comprising” claim ever written would be invalid for failing to describe every nuance of the claimed invention. Furthermore, the specification demonstrates to one skilled in the art that Appellants were in possession of the claimed genera of nucleic acid molecules.

(1) The Specification Reflects Appellants’ Possession of the Claimed Invention

The purpose of the written description requirement is to ensure that the inventor had possession of the claimed subject matter, *i.e.*, to ensure that the inventors actually invented what is claimed. *Gentry Gallery Inc. v. Berkline Corp.*, 134 F.3d 1473, 1479, 45 U.S.P.Q.2d 1498, 1503 (Fed. Cir. 1998); *Lockwood v. American Airlines*, 107 F.3d 1565, 1572, 41 U.S.P.Q.2d 1961, 1966 (Fed. Cir. 1997); *In re Alton*, 76 F.3d 1168, 1172, 37 U.S.P.Q.2d 1578, 1581 (Fed. Cir. 1996). If a person of ordinary skill in the art would, after reading the specification, understand that the inventors had possession of the claimed invention, even if not every nuance, then the written description has been met. *In re Alton*, 76 F.3d at 1175, 37 U.S.P.Q.2d at 1584. A person of ordinary skill in the art would, after reading the present specification, understand that Appellants had possession of SEQ ID NO: 48411 and complement thereof, as well as fragments thereof. Appellants have provided the nucleotide sequence required by the claims, *e.g.*, SEQ ID NO: 48411 and the complement thereof, and have provided fragments of the claimed sequence. Accordingly, Appellants have demonstrated possession of the claimed invention.

The fact that the claims at issue are intended to cover molecules that include fragments of the recited sequence, the recited sequence joined with additional sequences, or complements of the recited sequence, or nucleic acid molecules that share a claimed iden-

tity with the recited sequences, does not mean that Appellants were any less in possession of the claimed nucleic acid molecules.⁷ It is well-established law that use of the transitional term “comprising” properly leaves the claims “open for the inclusion of unspecified ingredients even in major amounts.” *Ex parte Davis*, 80 U.S.P.Q. 448, 450 (B.P.A.I. 1948); *Accord PPG Indus. v. Guardian Indus.*, 156 F.3d 1351, 1354, 48 U.S.P.Q.2d 1351, 1353-54 (Fed. Cir. 1998); *Moleculon Research Corp. v. CBS*, 793 F.2d 1261, 1271, 229 U.S.P.Q. 805, 812 (Fed. Cir. 1986).

The present application describes more than just the nucleotide sequence recited by the claims (SEQ ID NO: 48411). For example, the specification describes vectors comprising the claimed nucleic acid molecules (*see e.g.*, specification at page 23, line 28 through page 28, line 21) and describes how to make the nucleotide sequence and the libraries from which it was originally purified. *See, e.g.*, Example 1 at page 58, line 24 *et seq.* Furthermore, the addition of other nucleotides or detectable labels to the disclosed nucleotide sequences (*e.g.*, SEQ ID NO: 48411) is readily envisioned by one of ordinary skill in the art upon reading the present specification,⁸ as described for example at page 9, (describing sequences with labels to facilitate detection); as also described for example at page 19 (describing fusion peptide molecules encoded by the claimed nucleic acid molecules); and at page 51 (describing site-directed mutagenesis).

Moreover, the court determined, in *Enzo Biochem, Inc. v. Gen-Probe Inc.*, 296 F.3d 1316, 1321, 63 U.S.P.Q.2d 1609, 1610 (Fed. Cir. 2002), that the written description inquiry is a factual one determined on a case-by-case basis and that, in a given disclosure, “it may well be that various subsequences, mutations, and mixtures of those sequences

⁷ If the Examiner is arguing that no possession is shown because the precise claim language is not used in the specification, then it goes beyond what is required by the law. It is well-settled that the description of a claimed invention need not be *in ipsius verbis*. *Gentry Gallery v. Berkline Corp.*, 134 F.3d 1473, 1479, 45 U.S.P.Q.2d 1498, 1503 (Fed. Cir. 1998); *In re Alton*, 76 F.3d 1168, 1175, 37 U.S.P.Q.2d 1578, 1583 (Fed. Cir. 1996); *Martin v. Johnson*, 454 F.2d 746, 751, 172 U.S.P.Q. 391, 395 (C.C.P.A. 1972).

⁸ It is established patent jurisprudence that Appellant need not teach “conventional and well-known genetic engineering techniques.” *E.g., Ajinomoto Co. v. Archer-Daniels-Midland Co.*, 228 F.3d 1338, 1345, 56 U.S.P.Q.2d 1332, 1337 (Fed. Cir. 2000).

are also described to one of skill in the art.” *Enzo*, 296 F.3d at 1326-1327, 63 U.S.P.Q.2d at 1615. Furthermore, it is well established that claims “may be broader than the specific embodiment disclosed in a specification. *Ralston-Purina Co. v. Far-mor-Co*, 772 F.2d 1570, 1575, 227 U.S.P.Q. 177, 179 (Fed. Cir. 1985) (quoting *In re Rasmussen*, 650 F.2d 1212, 1215, 211 U.S.P.Q. 323, 326 (C.C.P.A. 1981).

(2) Appellants Have Described the Claimed Invention

The Examiner asserts that “the specification only provides sequences of the elected SEQ ID No:48411...”, and accordingly Applicants have allegedly not adequately disclosed the claimed genera of nucleic acid molecules. Final Action at page 4. As such, the Examiner appears to require that each nucleic acid molecule within the claimed genera must be described by its complete structure. Final Action at page 4. This requirement is totally unfounded. The Federal Circuit has elucidated a test for written description wherein a genus of nucleic acids may be described by a structural feature that distinguishes members of the claimed genus from non-members of the claimed genus. *Regents of the University of California v. Eli Lilly and Co.*, 119 F.3d 1559, 1568-69, 43 U.S.P.Q.2d 1398, 1406 (Fed. Cir. 1997). Appellants have satisfied that test for written description.

In particular, Appellants have disclosed common structural features, for example the nucleotide sequence of SEQ ID NO: 48411. For example, if a particular nucleic acid molecule contains the nucleotide sequence of SEQ ID NO: 48411, then it is a member of the claimed genus of nucleic acid molecules comprising a nucleic acid sequence of SEQ ID NO: 48411.⁹ Moreover, closely related nucleic acid molecules falling within the scope of the claimed invention are readily identifiable - they either contain the nucleic acid sequence of SEQ ID NO: 48411 (or complements or fragments thereof), or share a

⁹ The same argument applies with equal force to every genus of the claimed nucleic acid molecules. For example, if a nucleic acid molecule contains a nucleic acid sequence that has 95% identity with nucleotides 1 through 123 of SEQ ID NO: 48411, then it is a member of the claimed genus of nucleic acid molecules having between 90% and 100% identity with nucleotides 1 through 123 of SEQ ID NO: 48411. See claim 14.

claimed identity with SEQ ID NO: 48411 (or complements or fragments thereof), or they do not. The fact that the nucleic acid molecules may comprise additional sequences or variations is beside the point. Such modifications are readily envisioned by one of ordinary skill in the art and disclosed throughout the specification.

Moreover, if a particular nucleic acid molecule contains the claimed fragments of the nucleotide sequence of SEQ ID NO: 48411, then it is a member of the claimed genus of nucleic acid molecules comprising the recited fragments of a nucleic acid sequence of SEQ ID NO: 48411. Moreover, closely related nucleic acid molecules falling within the scope of the claimed invention are readily identifiable - they either contain a fragment of the recited fragment length of the nucleic acid sequence of SEQ ID NO: 48411 (or complements or fragments thereof), or share a claimed identity with SEQ ID NO: 48411 (or complements thereof), or they do not. The fact that the fragment nucleic acid molecules may comprise additional sequences is beside the point. Such modifications are readily envisioned by one of ordinary skill in the art and disclosed throughout the specification.

Thus, contrary to the Examiner's analysis, claims 11-15 are supported by an adequate written description pursuant to the requirements of 35 U.S.C. § 112, and the rejection should be reversed.

E. The Specification Provides An Adequate Written Description of the Claimed Invention: No New Matter Is Introduced

The adequacy of the written description of the claimed invention has been challenged by the Examiner because the inclusion of claim language "nucleotides 1 through 123 of SEQ ID NO: 48411" in claim 14 allegedly constitutes new matter. In order to comply with the written description requirement of 35 U.S.C. §112, Applicants must ensure that each portion of a claim is "expressly, implicitly, or inherently supported in the originally filed disclosure." MPEP §2163.05 at 2100-75; *Wertheim*, 541 F.2d 257, 191 U.S.P.Q. 90 (C.C.P.A. 1976). The analysis for numerical range limitations must take into account which ranges one skilled in the art would consider inherently supported by the original disclosure. *Id.* Nucleotides 1 through 123 are clearly present in Appellants' disclosure as filed. *See* SEQ ID NO: 48411 in the sequence listing. Additionally,

Appellants contemplate the use of fragment nucleic acid molecules throughout their disclosure. *See e.g.*, page 8 line 28 through page 9 line 2.

The present case is analogous to *In re Wertheim*, where the range 35%-60% was permitted when the original specification had described a range between 25% and 60%. By contrast, in *Wertheim*, the range at least 35% was deemed impermissible because it included percentages not originally disclosed, *i.e.*, those percentages greater than 60% may constitute new matter. In the present case, Appellants have described nucleotides 1 through 123 of SEQ ID NO: 48411 as well as fragments thereof. Furthermore, one of skill in the art can envision a nucleic acid molecule comprising a nucleic acid sequence having between 90% and 100% sequence identity with nucleotides 1 through 123 of SEQ ID NO: 48411 or a complete complement thereof based on Appellants' disclosure. *See e.g.*, specification at page 11, lines 5-20 and the sequence listing.

In contrast, the Examiner has not provided any support for the proposition that the claim limitation of "base pairs 1 through 123 of SEQ ID NO: 48411" is not described in Appellants' specification as originally filed. It is well-settled that the description of a claimed invention need not be in *ipsis verbis*. *Gentry Gallery v. Berkline Corp.*, 134 F.3d 1473, 1479, 45 U.S.P.Q.2d 1498, 1503 (Fed. Cir. 1998); *In re Alton*, 76 F.3d 1168, 1175, 37 U.S.P.Q.2d 1578, 1583 (Fed. Cir. 1996); *Martin v. Johnson*, 454 F.2d 746, 751, 172 U.S.P.Q. 391, 395 (C.C.P.A. 1972). Thus, the Examiner has not met the burden to impose a written description rejection of claim 14. ("[t]he Examiner has the initial burden of presenting by a preponderance of evidence why a person skilled in the art would not recognize in an applicant's disclosure a description of the invention defined by the claims.") *Wertheim*, 541 F.2d at 263, 191 U.S.P.Q. at 97, M.P.E.P. §2167.04 at 2100-73.

As such, written description of the claimed invention has been satisfied, and inclusion of claim language "nucleotides 1 through 123 of SEQ ID NO: 48411" in claim 14 does not constitute new matter. Appellants respectfully submit that the rejection of claim 14 under 35 U.S.C. §112, written description should be reversed.

F. The Claimed Nucleic Acid Molecules Are Novel

The novelty of the claimed invention has been challenged by the Examiner under 35 U.S.C. §102(b) because claim 13 is allegedly anticipated by Mahairas *et al.* ("Mahairas") (Accession No. AQ451805). "It is axiomatic that for prior art to anticipate under § 102 it has to meet every element of the claimed invention." *Hybritech Inc. v. Monoclonal Antibodies, Inc.*, 802 F.2d 1367, 231 U.S.P.Q. 81 (Fed. Cir. 1986). Further, "an anticipation rejection requires a showing that each limitation of a claim must be found in a single reference, practice, or device." *In re Donohue*, 766 F.2d 531, 226 U.S.P.Q. 619 (Fed. Cir. 1985).

In the Final Action, Claim 13 was erroneously rejected under 35 U.S.C. § 102(b) as allegedly being anticipated by Mahairas. The Examiner alleges that "absent a definition for the term 'fragment' of SEQ ID NO: 48411, one or more nucleotides are considered a fragment." Final Action at pages 6-7. However, this allegation fails to take account of the claim language, which recites a "fragment nucleic acid molecule having from about 30 to about 50 nucleotide residues, wherein said fragment nucleic acid molecule exhibits complete complementarity to a fragment of a second nucleic acid molecule having the nucleotide sequence of SEQ ID NO: 48411 or a complete complement thereof." The Examiner has not read the claims in light of Appellants' disclosure, as required, but rather has implied an interpretation, unsupported by evidence, that the claimed nucleic acid molecules encompass a fragment that "can be any nucleic acid fragment of about 30-50 bps long." Final Action at page 6. Such clearly unsupported conjecture is simply not a proper basis for an anticipation rejection.

The Examiner appears to suggest that because the nucleic acid molecule of Mahairas contains a fragment that is completely complementary to nucleotides 98-118 of SEQ ID NO: 48411, Mahairas is anticipatory. The Final Action alleges that "Mahairas *et al.* contains a fragment of around 30." Final Action at page 7. Such an interpretation of the phrase "about 30 to about 50 nucleotide residues" is not in accordance with the law. See *BJ Services Co. v. Halliburton Energy Services, Inc.*, 338 F.3d 1368, 67 U.S.P.Q.2d


1692 (Fed. Cir. 2003). From the decision in *BJ Services*, the term “about” in the present claims should be given its “plain and ordinary meaning.” *Id.* In *BJ Services*, the appellee attempted unsuccessfully to argue that 0.077 was “about 0.06.” However, according to *BJ Services*, 0.077 was deemed not to give “about 0.06” its plain and ordinary meaning. *See id.* The present case is analogous to *BJ Services*. The fragment of Mahairas as cited by the Examiner contains 21 nucleotides. As in *BJ Services*, this 21 nucleotide base pair fragment fails to give “about 30 nucleotides” its “plain and ordinary meaning.” *Id.*

Whatever else Mahairas may teach or suggest, it does not teach or suggest a substantially purified nucleic acid molecule comprising a fragment nucleic acid molecule having from about 30 to about 50 nucleotide residues, wherein said fragment nucleic acid molecule exhibits complete complementarity to a fragment of a second nucleic acid molecule having the nucleotide sequence of SEQ ID NO: 48411 or a complete complement thereof. The law requires that each and every element of a claimed invention is disclosed within a single prior art reference. *In re Bond*, 15 U.S.P.Q.2d 1566, 1567 (Fed. Cir. 1990). As such, the rejection of claim 13 as anticipated under 35 U.S.C. §102(b) by Mahairas is improper and should be reversed.

CONCLUSION

In view of the foregoing, it is respectfully requested that the Board of Patent Appeals and Interferences reverse the Rejections and that the subject application be allowed forthwith.

Respectfully submitted,



Date: December 22, 2006

Of Counsel

Lawrence M. Lavin, Jr. (Reg. No. 30,768)

Thomas E. Kelley (Reg. No. 29,938)

Monsanto Company

Thomas E. Holsten (Reg. No. 46,098)

David R. Marsh (Reg. No. 41,408)

ARNOLD & PORTER LLP

Attn: IP Docketing

555 Twelfth Street, NW

Washington, DC 20004-1206

202.942.5000 telephone

202.942.5999 facsimile

CLAIMS APPENDIX

11. A substantially purified nucleic acid molecule comprising a fragment nucleic acid molecule having from about 30 to about 50 nucleotide residues of a nucleic acid molecule having the nucleotide sequence of SEQ ID NO: 48411.

12. A substantially purified nucleic acid molecule comprising a fragment nucleic acid molecule having from about 50 to about 100 nucleotide residues of a nucleic acid molecule having the nucleotide sequence of SEQ ID NO: 48411.

13. A substantially purified nucleic acid molecule comprising a fragment nucleic acid molecule having from about 30 to about 50 nucleotide residues, wherein said fragment nucleic acid molecule exhibits complete complementarity to a fragment of a second nucleic acid molecule having the nucleotide sequence of SEQ ID NO: 48411 or a complete complement thereof.

14. A substantially purified nucleic acid molecule having between 90% and 100% sequence identity with nucleotides 1 through 123 of SEQ ID NO: 48411 or a complete complement thereof.

15. The substantially purified nucleic acid molecule of claim 14, wherein said substantially purified nucleic acid molecule has between 99% and 100% sequence identity with nucleotides 1 through 123 of SEQ ID NO: 48411 or a complete complement thereof.

16. A substantially purified nucleic acid molecule according to claim 15, wherein said nucleic acid molecule has the nucleic acid sequence of SEQ ID NO: 48411 or the complete complement thereof.

EVIDENCE APPENDIX

None

RELATED PROCEEDINGS APPENDIX

Attached are copies of Decisions on Appeal that issued in the following applications:

1. *In re Fisher*, 421 F.3d 1365, 76 U.S.P.Q.2d 1225 (Fed. Cir. 2005)
2. U.S. Appln. No. 09/619,643, BPAI Appeal No. 2002-2046
3. U.S. Appln. No. 09/654,617, BPAI Appeal No. 2003-1744
4. U.S. Appln. No. 09/620,392, BPAI Appeal No. 2003-1746
5. U.S. Appln. No. 09/540,232, BPAI Appeal No. 2003-1137
6. U.S. Appln. No. 09/440,687, BPAI Appeal No. 2003-1504
7. U.S. Appln. No. 09/565,240, BPAI Appeal No. 2003-1135
8. U.S. Appln. No. 09/540,215, BPAI Appeal No. 2003-0996
9. U.S. Appln. No. 09/552,087, BPAI Appeal No. 2004-1772
10. U.S. Appln. No. 09/206,040, BPAI Appeal No. 2002-0078



**IN THE UNITED STATES PATENT AND TRADEMARK OFFICE
BEFORE THE BOARD OF PATENT APPEALS AND INTERFERENCES**

In re Patent Application of:

Linda L. LUTFIYYA

Application Serial No.: 10/361,942

Filed: February 10, 2003

Confirmation No.: 8391

Art Unit: 1638

Examiner: Cynthia E. Collins

Attorney Docket No.: 16517.335

Title: Nucleic Acids and Other Molecules Associated with Transcription in Plants

Revised Appeal Brief under 37 C.F.R. § 41.37

Mail Stop Appeal Brief – Patents
Commissioner for Patents
P.O. Box 1450
Alexandria, VA 22313-1450

Sir:

This is an Appeal from the Final Rejection of claims in the above-captioned patent application. A Notice of Appeal was filed on August 3, 2006. Authorization to charge the official fees for this filing was given at the time of filing the original Appeal Brief on November 3, 2006. This Revised Appeal Brief is submitted in response to a Notification of Non-Compliant Appeal Brief dated November 16, 2006.

1. Real Party in Interest

The real party in interest is Monsanto Company, a Delaware corporation with offices at 800 North Lindbergh Boulevard, St. Louis, Missouri 63167.

2. Related Appeals and Interferences

The Appellant identified the following judicial proceeding, which may have a bearing on the Board's decision in the present Appeal. On May 27, 2004, the Real Party in Interest in the above-captioned matter filed an appeal to the United States Court of Appeals for the Federal Circuit ("Federal Circuit") from a decision by the Board in *In re Fisher*. (U.S. Patent Application. No. 09/619,643; BPAI Appeal No. 2002-2046; Fed. Cir. Case No. 04-1465). The Federal Circuit's decision in *In re Fisher* may have a bearing on the Board's decision with regard to at least one of the grounds of rejection in the present appeal. A copy of the Board's decision in Appeal No. 2002-2046, and a copy of *In re Fisher*, 421 F.3d 1365, 76 U.S.P.Q.2d 1225 (Fed. Cir. 2005) are attached hereto in the Related Proceedings Appendix.

In addition, Appellant also identifies the following additional Board decisions which may have a bearing on the instant appeal: U.S. Appln. No. 09/654,617, BPAI Appeal No. 2003-1744; U.S. Appln. No. 09/620,392, BPAI Appeal No. 2003-1746; U.S. Appln. No. 09/540,232, BPAI Appeal No. 2003-1137; U.S. Appln. No. 09/440,687, BPAI Appeal No. 2003-1504; U.S. Appln. No. 09/565,240, BPAI Appeal No. 2003-1135; U.S. Appln. No. 09/540,215, BPAI Appeal No. 2003-0996; U.S. Appln. No. 09/552,087, BPAI Appeal No. 2004-1772; and U.S. Appln. No. 09/206,040, BPAI Appeal No. 2002-0078. Copies of the Board's decisions in these Appeals are also attached hereto in the Related Proceedings Appendix.

Appellant also identifies the following pending appeals before the Board which may have a bearing on the instant appeal: U.S. Appln. No. 09/199,129, and U.S. Appln. No. 09/684,016.

3. Status of Claims

Claims 1, 8, 11, and 12 are pending. Claims 2 to 7, 9, and 10 were canceled without prejudice to, or disclaimer of, the subject matter claimed therein in the responses to the Office Action filed September 21, 2005 and August 3, 2006. Claims 1, 8, 11 and 12 stand finally rejected under 35 U.S.C. § 112, first paragraph and under 35 U.S.C. § 101. The Appellant appeals the rejections of claims 1, 8, 11, and 12.

4. Status of Amendments

On August 3, 2006, the Appellant filed an amendment subsequent to the final rejection concurrently with the Notice of Appeal, in which claims 9 and 10 were canceled. This amendment was entered for purposes of appeal. *See Advisory Action* mailed August 18, 2006.

A Second Amendment After Final Rejection was filed on November 3, 2006 amending claim 11 to correct its dependency. The Amendment is intended to clarify the claims on appeal.

5. Summary of Claimed Subject Matter

Independent Claim 1: The claimed subject matter of independent claim 1 is directed to an isolated nucleic acid molecule comprising a substantially purified nucleic acid molecule, or a complement thereof, comprising a nucleic acid sequence of SEQ ID NO: 1454. *Specification* at page 29, lines 12 to 13, and the Sequence Listing.

Independent Claim 8: The claimed subject matter of independent claim 8 is directed to a substantially purified nucleic acid molecule comprising a nucleic sequence, wherein the nucleic acid sequence shares between 100% and 98% and between 100% and 99% sequence identity with SEQ ID NO: 1454 or complement thereof. *Id.* at page 31, lines 5 to 18. SEQ ID NO: 1454

is shown in the specification to correlate to a known 14-3-3 transcription factor with an expectation value (E-value) of $1.2e-44$. *Id.* at Table 2.

A copy of the claims on appeal is attached hereto in the Claims Appendix.

6. Grounds of Rejection to be Reviewed on Appeal

The grounds of rejection to be reviewed in this Appeal are:

- (a) pending claims 11 and 12 stand rejected under 35 U.S.C. § 112, first paragraph, for allegedly failing to comply with the written description requirement;
- (b) pending claims 1, 8, 11, and 12 stand rejected under 35 U.S.C. § 112, first paragraph, for allegedly failing to comply with the enablement requirement; and
- (c) pending claims 1, 8, 11, and 12 stand rejected under 35 U.S.C. §§ 101 and 112 because the claimed invention is allegedly not supported by either a specific and substantial asserted utility or a well established utility.

7. Argument

A. Summary of Appellants' Position

The Appellant has provided an adequate description of the claimed nucleic acid molecules that demonstrates the Appellant's possession of the claimed invention. The claimed nucleic acid molecules, for example, the nucleic acid molecules comprising a nucleic acid sequence that shares between 100% and 98% sequence identity with SEQ ID NO: 1454, have been described by the recitation of common structural features, *e.g.*, nucleotide sequences of SEQ ID NO: 1454, which distinguish molecules in the claimed genus from molecules not in the claimed genus. Because the specification demonstrates that Appellant had possession of (and

has provided an adequate description of) the claimed genera of nucleic acid molecules, the specification satisfies the written description requirement of 35 U.S.C. § 112.

The Appellant has also provided sufficient disclosure in the specification to enable a person skilled in the art to make and/or use the invention. The specification clearly enables at least the methods of making and using the invention as set forth in the Examples. Furthermore, an analysis of the criteria presented by *In re Wands*, 858 F.2d 731, 8 U.S.P.Q.2d 1400 (Fed. Cir. 1998), indicates that no undue experimentation would be required to make and use the claimed invention. Thus, the specification satisfies the enablement requirement of 35 U.S.C. § 112.

Finally, the specification provides a specific and substantial utility for the claimed nucleic acid molecules - encoding 14-3-3 transcription factors, or fragments thereof. Appellant has provided a statistically significant correlation between the claimed nucleic acid sequences and known transcription factors. The utility of the known transcription factors is substantial and well-established; the correlation between the known transcription factors is specific to the claimed nucleic acid sequence. In setting forth a reasonable correlation between the known transcription factors and the claimed nucleic acid sequences, the Appellant has demonstrated sufficient utility of the claimed nucleic acid sequences to satisfy the requirements of 35 U.S.C. §§ 101 and 112.

B. The Specification Provides An Adequate Written Description of the Claimed Invention

The Examiner rejected pending claims 11 and 12 under 35 U.S.C. § 112, first paragraph, as allegedly “failing to comply with the written description requirement.” *Final Office Action* mailed May 3, 2006 at page 2.

Claims 11 and 12 are directed to, *inter alia*, nucleic acid sequences that share between 100% and 98% (claim 11) or between 100% and 99% (claim 12) sequence identity with SEQ ID NO: 1454 or its complement. The Appellant has disclosed the nucleic acid sequence of SEQ ID NO: 1454. *See, e.g., Sequence Listing*. However, the Examiner argues that the Appellant has allegedly not described “a representative number of species falling within the scope of the claimed genus which encompasses numerous undisclosed and uncharacterized variants of SEQ ID NO: 1454, nor the structural features unique to the genus that are correlated with a common function.” *Final Office Action* at page 3. Contrary to the Examiner’s allegations, however, the specification provides an adequate description of the claimed invention because it demonstrates to one skilled in the art that the Appellant was in possession of the claimed genera of nucleic acid molecules when the application was filed.

The purpose of the written description requirement is to ensure that the inventor had possession of the claimed subject matter, *i.e.*, to ensure that the inventor actually invented what is claimed. *Gentry Gallery Inc. v. Berkline Corp.*, 134 F.3d 1473, 1479, 45 U.S.P.Q.2d 1498, 1503 (Fed. Cir. 1998); *Lockwood v. American Airlines*, 107 F.3d 1565, 1572, 41 U.S.P.Q.2d 1961, 1966 (Fed. Cir. 1997); *In re Alton*, 76 F.3d 1168, 1172, 37 U.S.P.Q.2d 1578, 1581 (Fed. Cir. 1996). In accordance with this purpose, Appellant need not “describe,” in the sense of Section 112, all things that are encompassed by the claims. To contend otherwise would contradict established jurisprudence, which teaches that a patent may be infringed by technology developed after a patent issues. *United States Steel Corp. v. Phillips Petroleum Co.*, 865 F.2d 1247, 1251, 9 U.S.P.Q.2d 1461, 1464 (Fed. Cir. 1989). A related, and equally well-established principle of patent law is that claims “may be broader than the specific embodiment disclosed in

a specification.” *Ralston-Purina Co. v. Far-mor-Co*, 772 F.2d 1570, 1575, 227 U.S.P.Q. 177, 179 (Fed. Cir. 1985) (quoting *In re Rasmussen*, 650 F.2d 1212, 1215, 211 U.S.P.Q. 323, 326 (C.C.P.A. 1981)). Thus, in order for the Appellant to describe each and every molecule encompassed by the claims, it is not required that every aspect of those nucleic acid molecules (e.g., “essential regions”) be disclosed. *In re Alton*, 76 F.3d at 1175, 37 U.S.P.Q.2d at 1584 (if a person of ordinary skill in the art would, after reading the specification, understand that the inventors had possession of the claimed invention, even if not every nuance, then the written description has been met).

It is well-settled law that each nucleic acid molecule within a claimed genus does not need to be described by its complete structure. An adequate written description of a genus of nucleic acids, such as those recited in claims 11 and 12, may be achieved by either “a recitation of a representative number of [nucleic acid molecules], defined by nucleotide sequence, falling within the scope of genus or of a recitation of structural features common to the members of the genus.” *Regents of the University of California v. Eli Lilly and Co.*, 119 F.3d 1559, 1568-69, 43 U.S.P.Q.2d 1398, 1406 (Fed. Cir. 1997) (emphasis added). The feature relied upon to describe the claimed genus must be capable of distinguishing members of the claimed genus from non-members. *Id.* In contrast to the mere name “cDNA” provided in *Eli Lilly*, the Appellant has provided a detailed chemical structure by way of the nucleic acid sequence of SEQ ID NO: 1454, as well as complements and specified variations thereof. This chemical structure clearly distinguishes molecules in the claimed genus from molecules not in the claimed genus. The Appellant has therefore satisfied the *Eli Lilly* test for written description.

The fundamental factual inquiry for satisfying the written description requirement is whether the specification conveys with reasonable clarity to those skilled in the art, as of the filing date sought, that the Appellant was in possession of the invention as now claimed. *See, e.g., Vas-Cath, Inc. v. Mahurkar*, 935 F.2d 1555, 1563-64, 19 U.S.P.Q. 2d 1111, 1117 (Fed. Cir. 1991). An applicant shows possession of the claimed invention by describing the claimed invention with all of its limitations. *Lockwood v. American Airlines, Inc.*, 107 F.3d 1565, 1572, 41 U.S.P.Q.2d 1961, 1966 (Fed. Cir. 1997); M.P.E.P. § 2163.02. In light of the disclosure made by the Appellant in the specification one of ordinary skill in the art at the time the application was filed would have readily recognized that the Appellant was in possession of the invention as claimed. The Examiner has offered no evidence to demonstrate why one of ordinary skill in the art would reasonably doubt that the invention has not been adequately described in the present disclosure. Therefore, the Examiner has not met the burden to impose a written description rejection.

The Appellant has provided a detailed chemical structure, *i.e.*, the nucleic acid sequence, of the claimed SEQ ID NO: 1454. Nucleic acid molecules falling within the scope of claims 11 and 12 are readily identifiable – they share between 100% to 98% identity to SEQ ID NO: 1454 or its complement or they do not. One of ordinary skill in the art can readily identify whether a particular sequence meets these characteristics. The fact that the nucleic acid molecules may comprise additional sequences or variations is beside the point. Such modifications are readily envisioned by one of ordinary skill in the art and disclosed throughout the specification.

In conclusion, the Appellant respectfully submits that claims 11 and 12 are fully supported by an adequate written description pursuant to the requirements of 35 U.S.C. § 112.

Therefore, the Appellant respectfully requests that the Board reverse the written description rejection of claims 11 and 12 under 35 U.S.C. § 112, first paragraph.

C. The Claimed Nucleic Acids are Fully Enabled by the Specification

The Examiner rejected pending claims 1, 8, 11, and 12 under 35 U.S.C. § 112, first paragraph, as allegedly “failing to comply with the enablement requirement.” *Final Office Action* at page 3. The Examiner states that the enablement rejection is “predicated on the quantum of experimentation required to determine how a nucleic acid of sequence of SEQ ID NO: 1454 may be used.” *Id.* at page 6. The Examiner suggests that “[s]uch guidance is necessary because polypeptide function cannot be reliably predicted on the basis of amino acid sequence homology alone, because polypeptide function cannot be reliably predicted on the basis of amino acid sequence homology to 14-3-3 proteins, and because polypeptide function cannot be reliably predicted on the basis of a partial amino acid sequence.” *Id.* In other words, the Examiner appears to be asserting that the specification does not enable claims 1, 8, 11, and 12 because one of ordinary skill in the art would require undue experimentation to use the claimed invention.

The Examiner relies on several publications to support the assertion that sequence homology alone does not reliably predict the function of an encoded polypeptide. *Id.* at page 10. However, the references cited by the Examiner support the identification of protein function using sequence homology. As Whisstock *et al.* acknowledge, sequence homology-based “methods provide reasonable guesses at function....” *See, e.g., Whisstock et al. Quarterly Reviews of Biophysics*, 36, 3: 307-340 (2003) (Abstract). Indeed the references cited by the Examiner also show that one of ordinary skill in the art is well-aware of 14-3-3 transcription factors, the numerous cellular roles that such transcription factors play and methods for

determining 14-3-3 transcription factor binding. For example, Finnie, C. *et al.* ("14-3-3 Proteins: Eukaryotic Regulatory Proteins With Many Functions", *Plant Mol. Biol.* **40**(4), 545–554, 1999) provide a summary of some of the knowledge the skilled artisan possesses of 14-3-3 transcription factor family receptors and their enzymatic roles in plant cells. Finnie *et al.* review studies investigating 14-3-3 transcription factors and assays for categorizing such factors, for example by phosphorylation of 14-3-3 binding motifs. It is well established that in the context of enablement a patent application "need not teach, and preferably omits, what is well known in the art." *Hybritech, Inc. v. Monoclonal Antibodies, Inc.*, 802 F.2d 1367, 1384 (Fed. Cir. 1986) citation omitted. The teachings of the present specification taken in combination with the knowledge of those skilled in the art provide adequate guidance regarding the identification of 14-3-3 transcription factor proteins.

Moreover, an analysis of the criteria presented in *In re Wands* supports the Appellant's position that no undue experimentation would be required to make and use the claimed invention. *See In re Wands*, 858 F.2d 731, 737, 8 U.S.P.Q.2d 1400, 1404 (Fed. Cir. 1998). The Appellant has provided considerable direction and guidance, and has presented working examples such that it is well within the level of ordinary skill in the art to practice the claimed invention without undue experimentation. For example, the specification discusses nucleic acid molecules isolated from various *Arabidopsis thaliana* tissue, as well as homologous sequences. *See Specification*, page 28, line 2 through page 32, line 18 and page 81, line 11 through page 84, line 19. In addition, the specification discusses the preparation of constructs using the claimed sequences. *See Specification*, page 40, line 4 through page 48, line 2, and page 82, lines 18-22. The specification also discusses that the nucleic acid molecules can encode transcription factors

or fragments thereof. *See Specification*, page 32, line 20 through page 37, line 15. Moreover, the specification provides sequence homology statistics with regard to the disclosed sequences and further discloses the functions of various transcription factors, including 14-3-3 transcription factors. *See Specification*, page 13, line 1 through page 28, line 1 (Table 1) and Table 2. Taken in combination, such disclosure provides adequate direction to those skilled in the art of how to make and use the claimed invention as currently claimed.

The first *Wands* criterion is the quantity of experimentation necessary. The “make-and-test” quantum of experimentation is reduced by the extensive knowledge, *e.g.*, of conservative nucleotide substitutions, expression systems, and enzyme assay conditions, to which a person of ordinary skill in the art has access. One of ordinary skill in the art is sufficiently guided by the Appellant’s disclosure, which sets forth nucleic acid molecules as well as the fragments or complements thereof. Further, as stated above, practitioners in the art are guided by the high level of skill in the art and the present disclosure of the specification (*see, e.g., Specification* at page 13 (Table 1)). Performing routine and well-known steps, such as sequence alignment protocols, molecular weight determination, and known enzyme assays, cannot create undue experimentation even if it is laborious. *See In re Angstadt*, 537 F.2d 498, 504, 190 U.S.P.Q. 214, 218-219 (C.C.P.A. 1976).

The second and third *Wands* criteria relate to the amount of direction or guidance given, and the presence or absence of working examples. Again, the specification provides evidence of sequence identity via expectation values, discloses enzyme function, and discusses the use of the claimed SEQ ID NOs to isolate additional sequences within a genome. *See, Specification* at page 14 (Table 1), Examples 1 and 2, the Sequence Listing and Table 2. Based on such

disclosure, one of ordinary skill in the art would be enabled to make and use the invention commensurate in scope with the claims.

The fourth, fifth, and sixth *Wands* criteria focus on the nature of the invention, the state of the art, and the relative skill in the art. The specification provides a detailed description of the nucleic acid sequences required by the claims, and further describes amino acid sequences derived therefrom, and constructs and methods of use related thereto. *Id.* at page 28, line 3 through page 48, line 2, and the Sequence Listing (describing polypeptide molecules encoded by the nucleic acid sequences of the present invention, homologues and other modifications, and methods of producing or expressing peptides or fragments of peptides), and page 48, line 3 through page 59, line 20 (describing use of the claimed nucleic acid molecules in methods of transforming plants). Practitioners in this art are guided by considerable knowledge and resources on the conditions and approaches that can be utilized to identify, confirm, and introduce into other hosts, nucleic acid and amino acid sequences.

The seventh criterion considers the predictability of the art. The Appellant respectfully asserts, as discussed *infra*, that the specification discloses sufficient guidance to render the results of transformations with the claimed nucleic acid molecules predictable. *Id.* at page 28, line 3 through page 48, line 2, and Table 2. Furthermore, the specification provides sufficient guidance to one of skill in the art to make and use the claimed nucleic acid molecules. *Id.* at page 14 of Table 1 (describing 14-3-3 proteins), and page 65, line 1 through page 67, line 10 (citing methods for assaying gene expression).

The eighth criterion focuses on the breadth of the claims. Enablement is satisfied when the disclosure “adequately guide[s] the art worker to determine, without undue experimentation,

which species among all those encompassed by the claimed genus possess the disclosed utility”.
See In re Vaeck, 947 F.2d 488, 496, 20 U.S.P.Q.2d 1438, 1445 (Fed. Cir. 1991). In the present case, one of skill in the art is specifically guided by the disclosure to look to, *e.g.*, sequence identity data in making that determination.

In addition, the Examiner has not met the evidentiary burden to impose an enablement rejection. A specification that discloses how to use a claimed invention “must be taken as in compliance with the enabling requirement of the first paragraph of § 112 unless there is reason to doubt the objective truth of the statements contained therein.” *In re Brana*, 51 F.3d 1560, 1566, 34 U.S.P.Q.2d 1436, 1441 (Fed. Cir. 1995), *quoting In re Marzocchi*, 439 F.2d 220, 223, 169 U.S.P.Q. 367, 369 (C.C.P.A. 1971) (emphasis in original). The Examiner has provided neither evidence supporting the rejection nor any explanation of why the specification allegedly fails to enable the claimed nucleic acid molecules. *See In re Wright*, 999 F.2d 1557, 1561-62, 27 U.S.P.Q.2d 1510, 1513 (Fed. Cir. 1993); *Ex parte Lemak*, 210 U.S.P.Q. 306, 307 (B.P.A.I. 1981) (“pure conjecture” does not substantiate rejection for lack of enablement).

In conclusion, the Appellant respectfully submits that the above analysis illustrates that claims 1, 8, 11, and 12 are fully enabled by the specification. *Cf. Johns Hopkins University v. CellPro*, 152 F.3d 1342, 1361, 47 U.S.P.Q.2d 1705, 1719 (Fed. Cir. 1998) (“the enablement requirement is met if the description enables any mode of making and using the invention”) (emphasis added), *quoting Engel Indus. v. Lockformer Co.*, 946 F.2d 1528, 1533, 20 U.S.P.Q.2d 1300, 1304 (Fed. Cir. 1991). Therefore, the Appellant respectfully requests reversal of the enablement rejection of claims 1, 8, 11, and 12 under 35 U.S.C. § 112, first paragraph.

D. The Claimed Nucleic Acids Have Utility

The Appellant has provided nucleic acid sequences that the specification shows are correlated to underlying genes of a known function. One of the utilities of SEQ ID NO: 1454 is to encode an enzyme or complement thereof that is classified as a member of the 14-3-3 transcription factor family. *Specification* at page 14. The specification discloses that the claimed nucleic acid molecule correlates to a known 14-3-3 transcription factor with an expectation value (E value) of $1.2e-44$, suggesting a statistically significant homology to known 14-3-3 transcription factors. *Id.* at Table 2. Such use provides a specific and substantial utility.

The Examiner's position appears to be that the claimed invention lacks utility "because polypeptide function cannot be reliably predicted on the basis of amino acid sequence homology alone." *Final Office Action* at page 13. The Examiner cites Whisstock, J.C. *et al.* ("Prediction of Protein Function from Protein Sequence and Structure," *Q. Rev. Biophys.* **36**(3), 307–340, 2003), Emi, T. ("Specific Binding of vfl4-3-3a Isoform to the Plasma Membrane H^+ -ATPase in Response to Blue Light and Fusicoccin in Guard Cells of Broad Bean", *Plant Physiol.* **125**(2), 1115–1125, 2001) and Finnie, C. *et al.* as evidence in support.

Although the references cited by the Office may generally describe some level of unpredictability of the relationship between sequence/structure and function, the Examiner has provided no support for the allegation that use of the claimed nucleic acid molecules to encode 14-3-3 transcription factors, or fragments thereof, lacks specificity. Such a utility is not vague or unknown. In fact, the references that the Examiner has cited acknowledge the important roles of 14-3-3 transcription factors. The asserted utility, to encode a 14-3-3 transcription factor or fragment thereof is a well-defined use that not any nucleic acid sequence can perform.

An Examiner must accept a utility by an Applicant unless the Office has evidence or sound scientific reasoning to rebut the assertion. *See, In re Oetiker*, 977 F.2d 1443, 1445, 24 U.S.P.Q.2d 1443, 1444 (Fed. Cir. 1992), emphasis added. The Office has acknowledged that sequence homology is sufficient to establish utility. *See, e.g., Utility Examination Guidelines*, 66 Fed. Reg. 1092 (Jan. 5, 2001), as incorporated into the Manual of Patent Examining Procedure (“MPEP”), at § 2107; *See also*, MPEP § 2107.03 II. The Office’s Utility Guidelines provide that “when a patent application claiming a nucleic acid asserts a specific, substantial, and credible utility, and bases the assertion upon homology to existing nucleic acids or proteins having an accepted utility, the asserted utility must be accepted by the examiner unless the Office has sufficient evidence or sound scientific reasoning to rebut such as assertion.” Federal Register 66(4):1096, emphasis added. The Examiner has neither provided sufficient evidence nor sound scientific reasoning. What the Examiner has provided are a few scientific publications indicating that sequence and structural homology may not always rigorously be correlated with functionality.

Moreover, “a ‘rigorous correlation’ need not be shown in order to establish practical utility; ‘reasonable correlation’ is sufficient.” *See, Fujikawa v. Wattanasin*, 93 F.3d 1559, 1565, 39 U.S.P.Q.2d 1895, 1900 (Fed. Cir. 1996), emphasis added. “An Applicant can establish this reasonable correlation by relying on statistically relevant data documenting the activity of the compound or composition, arguments or reasoning, documentary evidence, or any combination thereof.” M.P.E.P. § 2107.03, at page 2100-43. The specification provides such a reasonable correlation through the statistically significant homology of SEQ ID NO: 1454 with a known 14-3-3 transcription factor (E value = 1.2e-44). Indeed one of the references cited by the Examiner

(Whisstock *et al.*) states that sequence homology “methods provide reasonable guesses at function.” Whisstock *et al.* Abstract, emphasis added. A reasonable correlation between the claimed nucleic acid sequence and a known protein is all that is legally required, and the Appellant has thoroughly satisfied this requirement.

The specification provides extensive evidence based on sequence identity that the claimed nucleic acid molecules encode a polypeptide having sequence homology to a 14-3-3 transcription factor with an e-value of 1.2e-44. *See Specification* at Table 2. The specification also indicates by way of the description of the enzymatic function of 14-3-3 transcription factors that the specified enzyme has well-known enzymatic function in the art. *See Specification* at page 14, Table 1. Further 14-3-3 transcription factors and uses thereof were known in the art, as suggested by the references cited by the Examiner.

The Examiner provides no support to show that SEQ ID NO: 1454 does not function as described by the specification. The Examiner cannot merely cite general scientific publications that show that in some instances sequence and structural homology does not rigorously correlate with functionality. These references do not rise to the threshold of sound scientific reasoning necessary to rebut the Appellant’s assertion of utility as required by the Utility Guidelines. The Examiner also cannot dismiss the holding of *Fujikawa* by stating that the compound at the center of the dispute in *Fujikawa* is different from the present application. *Final Office Action* at page 16. The fact that the chemical compound in *Fujikawa* is different from that in the present appeal does not ablate the legal significance of *Fujikawa*. Moreover, because *Fujikawa* has not been overruled, or its legal weight otherwise diminished, the Examiner is bound by its holding.

The claimed nucleic acid molecules have been asserted to encode a 14-3-3 transcription factor or fragment thereof. The specification provides ample correlation between the claimed nucleic acid molecule and 14-3-3 transcription factor proteins. Accordingly, the assertion of the use of the claimed nucleic acid molecules to encode a 14-3-3 transcription factor or fragment thereof satisfies the utility requirement of 35 U.S.C. § 101.

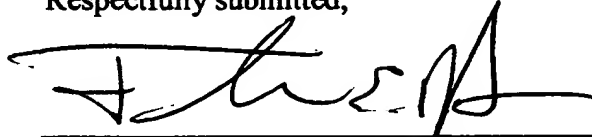
In addition to encoding a 14-3-3 transcription factor or fragment thereof, the specification describes multiple other utilities for the present invention that are independent of the sequence's ability to encode a 14-3-3 transcription factor or fragment thereof, including use as a molecular marker, obtain promoters, and use as probes for assisting in the isolation of full-length cDNAs or genes, isolation of homologous sequences, and the detection of gene expression. *See Specification* at page 61, line 14 *et seq*, under the heading "Exemplary Uses." Any of these utilities described alone is enough to satisfy 35 U.S.C. § 101. These uses are specific and substantial because, for example, the claimed nucleic acid molecules can be used as a marker or probe to the 14-3-3 transcription factor. Because the Appellant need only establish a single utility to satisfy 35 U.S.C. § 101, and has done so with reasonable correlation in the present case, the premise of the rejection under 35 U.S.C. § 101 is incorrect. Moreover, because the claimed nucleic acid sequences have utility under 35 U.S.C. § 101, they also have utility under 35 U.S.C. § 112, first paragraph.

In conclusion, the Appellant respectfully requests that the Board reverse the rejection of claims 1, 8, 11, and 12 under 35 U.S.C. §§ 101 and 112.

CONCLUSION

In view of the foregoing, the Appellant respectfully requests that the Board of Patent Appeals and Interferences reverse the Rejections and that the subject application be allowed forthwith.

Respectfully submitted,



Thomas E. Holsten (Reg. Atty. No. 46,098)
Gautam Prakash, Ph.D. (Reg. Agent No. 53,481)
David R. Marsh (Reg. Atty. No. 41,408)

Date: December 18, 2006

Of Counsel:

Lawrence M. Lavin, Jr. (Reg. No. 30,768)
Thomas E. Kelley (Reg. No. 29,938)
Monsanto Company

Arnold & Porter LLP
555 Twelfth Street, N.W.
Attn: IP Docketing
Washington, DC 20004

Tel: 202-942-5000
Fax: 202-942-5999

CLAIMS APPENDIX

(Claims as Pending)

Claim 1. A substantially purified nucleic acid molecule comprising a nucleic acid sequence of SEQ ID NO: 1454.

Claims 2 to 7. (Canceled)

Claim 8. A substantially purified nucleic acid molecule consisting of a nucleic acid sequence of SEQ ID NO:1454 or complement thereof.

Claims 9 and 10. (Canceled)

Claim 11. The substantially purified nucleic acid molecule of claim 10, wherein said nucleic acid sequence shares between 100% and 98% sequence identity with SEQ ID NO: 1454 or complement thereof.

Claim 12. The substantially purified nucleic acid molecule of claim 11, wherein said nucleic acid sequence shares between 100% and 99% sequence identity with SEQ ID NO: 1454 or complement thereof.

CLAIMS APPENDIX

(Claims as Amended)

Claim 1. A substantially purified nucleic acid molecule comprising a nucleic acid sequence of SEQ ID NO: 1454.

Claims 2 to 7. (Canceled)

Claim 8. A substantially purified nucleic acid molecule consisting of a nucleic acid sequence of SEQ ID NO:1454 or complement thereof.

Claims 9 and 10. (Canceled)

Claim 11. A substantially purified nucleic acid molecule wherein said nucleic acid molecule comprises a nucleic acid sequence that shares between 100% and 98% sequence identity with SEQ ID NO: 1454 or complement thereof.

Claim 12. The substantially purified nucleic acid molecule of claim 11, wherein said nucleic acid sequence shares between 100% and 99% sequence identity with SEQ ID NO: 1454 or complement thereof.

EVIDENCE APPENDIX

None

RELATED PROCEEDINGS APPENDIX

Attached are copies of Decisions on Appeal that issued in the following applications:

1. *In re Fisher*, 421 F.3d 1365, 76 U.S.P.Q.2d 1225 (Fed. Cir. 2005)
2. U.S. Appln. No. 09/619,643, BPAI Appeal No. 2002-2046
3. U.S. Appln. No. 09/654,617, BPAI Appeal No. 2003-1744
4. U.S. Appln. No. 09/620,392, BPAI Appeal No. 2003-1746
5. U.S. Appln. No. 09/540,232, BPAI Appeal No. 2003-1137
6. U.S. Appln. No. 09/440,687, BPAI Appeal No. 2003-1504
7. U.S. Appln. No. 09/565,240, BPAI Appeal No. 2003-1135
8. U.S. Appln. No. 09/540,215, BPAI Appeal No. 2003-0996
9. U.S. Appln. No. 09/552,087, BPAI Appeal No. 2004-1772
10. U.S. Appln. No. 09/206,040, BPAI Appeal No. 2002-0078



IN THE UNITED STATES PATENT AND TRADEMARK OFFICE
BEFORE THE BOARD OF PATENT APPEALS AND INTERFERENCES

In re application of:

Joseph R. BYRUM *et al.*

Appln. No.: 09/199,129

Filed: November 24, 1998

Title: Nucleic Acid Molecules and Other Molecules Associated with Plants

Confirmation No.: 3322

Art Unit: 1633

Examiner: Janet L. EPPS FORD

Atty. Docket: 16517.140

APPELLANTS' AMENDED BRIEF

Mail Stop Appeal Brief – Patents

Commissioner for Patents

P.O. Box 1450

Alexandria, Virginia 22313-1450

Sir:

This is an Appeal from the Final Rejection of all claims pending in the above-captioned patent application. A Notice of Appeal was filed on August 31, 2005. Authorization to charge the official fees for this filing is given in the accompanying transmittal letter.

1. Real Party in Interest

The real party in interest is Monsanto Company, a Delaware corporation with offices at 800 North Lindbergh Boulevard, St. Louis, Missouri 63167.

2. Related Appeals and Interferences

Appellant identifies the following judicial proceeding, which may have a bearing on the Board's decision in the present Appeal. On May 27, 2004, the Real Party in Interest in the above-captioned matter filed an appeal to the United States Court of Appeals for the Federal Circuit ("Federal Circuit") from a decision by the Board in *In re Fisher*. (U.S. Appln. No. 09/619,643, BPAI Appeal No. 2002-2046, Fed. Cir. Case No. 04-1465). The Federal Circuit's decision in *In re Fisher* may have a bearing on the Board's decision with regard to at least one of the grounds of rejection in the present appeal. A copy of the

Board's decision in Appeal No. 2002-2046, and a copy of *In re Fisher*, 412 F.3d 1365 (Fed. Cir. 2005) are attached hereto in the Related Proceedings Appendix.¹

In addition, Appellant also identifies the following additional Board decisions which may have a bearing on the instant appeal: U.S. Appln. No. 09/654,617, BPAI Appeal No. 2003-1744; U.S. Appln. No. 09/620,392, BPAI Appeal No. 2003-1746; U.S. Appln. No. 09/540,232, BPAI Appeal No. 2003-1137; U.S. Appln. No. 09/440,687, BPAI Appeal No. 2003-1504; U.S. Appln. No. 09/565,240, BPAI Appeal No. 2003-1135; U.S. Appln. No. 09/540,215, BPAI Appeal No. 2003-0996; U.S. Appln. No. 09/552,087, BPAI Appeal No. 2004-1772; and U.S. Appln. No. 09/206,040, BPAI Appeal No. 2002-0078. Copies of the Board's decisions in these Appeals are also attached hereto in the Related Proceedings Appendix.²

Appellant also identifies the following pending appeals before the Board which may have a bearing on the instant appeal: U.S. Appln. No. 09/233,218, BPAI Appeal No. 2004-1725; U.S. Appln. No. 09/540,234, BPAI Appeal No. 2003-1073; U.S. Appln. No. 09/333,535, BPAI Appeal No. 2003-1939; U.S. Appln. No. 09/666,355, BPAI Appeal No. 2004-1034; U.S. Appln. No. 09/552,086, BPAI Appeal No. 2003-1074; U.S. Appln. No. 09/637,086, BPAI Appeal No. 2004-1273; U.S. Appln. No. 09/540,235, BPAI Appeal No. 2004-1275; U.S. Appln. No. 09/553,094, BPAI Appeal No. 2004-1406; U.S. Appln. No. 09/267,199, BPAI Appeal No. 2004-2136; U.S. Appln. No. 09/521,640, BPAI Appeal No. 2004-1666; U.S. Appln. No. 09/371,146, BPAI Appeal No. 2004-1272; U.S. Appln. No. 09/421,106, BPAI Appeal No. 2004-1773; and U.S. Appln. No. 09/732,627, BPAI Appeal No. 2004-1480.³

¹ Copies of the Board's decision in Appeal No. 2002-2046, and a copy of *In re Fisher*, 412 F.3d 1365 (Fed. Cir. 2005) were submitted with Appellants' Brief filed October 31, 2005; however, for the Board's convenience, additional copies are submitted herewith.

² Copies of the Board's decisions in these appeals were submitted with Appellants' Brief filed October 31, 2005; however, for the Board's convenience, additional copies are submitted herewith.

³ At the time Appellant filed Appellants' Brief filed October 31, 2005, it had been requested that these appeals be suspended. Those appeals were subsequently suspended and were withdrawn from the Board of Patent Appeals and Interferences. None of these appeals are currently pending.

3. Status of Claims

Claims 4-12 are pending. Claims 2, 3 and 13-17 were cancelled without prejudice to or disclaimer of the subject matter claimed therein in a amendment filed July 2, 2002. Claim 1 was cancelled without prejudice to or disclaimer of the underlying subject matter in an amendment filed January 26, 2005. Claims 4-12 stand finally rejected under 35 U.S.C. §§101 and 112, first paragraph. Appellant appeals all of the rejections of claims 4-12.

4. Status of Amendments

Appellant has not filed any responses subsequent to Final Rejection in this case.

5. Summary of the Claimed Subject Matter

The claimed subject matter is directed to a transformed plant having a nucleic acid molecule which comprises: (a) an exogenous promoter region which functions in a plant cell to cause the production of an mRNA molecule; (b) a structural nucleic acid molecule comprising a nucleic acid sequence of SEQ ID NO: 1 and complement thereof; and (c) a 3' non-translated sequence that functions to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of said mRNA molecule. Specification at page 12, lines 7-15, page 57, line 13 through page 62, line 13, page 19, line 11 through page 24, line 2, and page 62, lines 14-19. The claimed subject matter is also directed to a method of determining a level or pattern in a plant cell or plant tissue of a protein in a plant comprising: (a) incubating, under conditions permitting nucleic acid hybridization, a marker nucleic acid molecule which specifically hybridizes to a nucleic acid molecule having the nucleic acid sequence of SEQ ID NO: 1 or complement thereof, with a complementary nucleic acid molecule obtained from the plant cell or plant tissue, where hybridization permits the detection of an mRNA for the protein; (b) permitting hybridization between said marker nucleic acid molecule and said complementary nucleic acid molecule obtained from said plant cell or plant tissue; and (c) detecting the level or pattern of said complementary nucleic acid, where the detection of said complementary nucleic acid is predictive of the level or pattern of said protein. Specification at page 15, lines 3-16, specification at page 20, line 17 through page 22, line 12, page 33, line 19 through page 34, line 20.

6. Grounds of Rejection to be Reviewed on Appeal

The grounds of rejection to be reviewed in this Appeal are:

(a) pending claims 4-12 stand rejected under 35 U.S.C. § 101, for allegedly not being supported by a specific asserted utility or a well established utility;

(b) pending claims 4-12 stand rejected under 35 U.S.C. § 112, first paragraph for alleged lack of enablement because the claimed invention purportedly lacks utility; and

(c) pending claims 4-12 stand rejected under 35 U.S.C. § 112, first paragraph for alleged insufficiency of written description.

A. Grouping of Claims

Claims 4-12 are pending in this application. All of the claims at issue do not stand or fall together. The separate patentability of claims 4-7 and 8-12 is addressed together in Sections 7.B.(1)(a) and 7.B.(1)(b) below. In addition, the separate patentability of claim 9, and the separate patentability of claim 10 are addressed in Sections 7.B.(1)(c) and 7.B.(1)(d), respectively.

7. Argument

A. Summary of Appellants' Position

As the Supreme Court said in *Brenner v. Manson*, the “basic quid pro quo contemplated by the Constitution and the Congress for granting a patent monopoly is the benefit derived by the public from an invention with substantial utility ... where specific benefit exists in currently available form.” 383 U.S. 519, 534-35, 148 U.S.P.Q. 689, 695 (1966). Applicants have met their part of the bargain – they have disclosed transformed plants and methods that, in their current form, provide at least one specific benefit to the public, for example, use of the transformed plants in a breeding program. This benefit is specific, and it is a “real world” or substantial benefit. Because the claimed plants and methods provide at least this benefit, they satisfy the utility requirement of 35 U.S.C. § 101. Because the specification teaches how to make and use the claimed plants and methods for the disclosed utilities, the enablement requirement of 35 U.S.C. § 112 has likewise been met.

Furthermore, Applicants have provided an adequate description of the plants and methods that demonstrates Applicants' possession of the claimed invention. Each genus of

nucleic acid molecule in the claimed plants and methods, *e.g.*, transformed plants comprising the nucleic acid molecules comprising a nucleic acid sequence of SEQ ID NO: 1, for example, has been described by the recitation of a common structural feature – the nucleotide sequence of SEQ ID NO: 1 – which distinguishes plants having molecules within the genus from plants having molecules outside of the genus. Because the specification demonstrates that Applicants have possession of (and have provided an adequate description of) the claimed plants and methods, the specification satisfies the written description requirement of 35 U.S.C. § 112.

B. The Claimed Plants and Methods Have Legal Utility

Pending claims 4-12 were erroneously rejected under 35 U.S.C. § 101 because the claimed invention was allegedly “not supported by either a specific and/or substantial utility or a well-established utility.” Final Action at pages 2-6. According to the Final Action, the “specification does not disclose or provide any evidence that points to a specific or substantial biologically significant activity for nucleic acid comprising SEQ ID NO: 1, a plant comprising said nucleic acid, or a method of identifying an unknown protein, such that another non-asserted utility would be well-established.” *Id.* at page 4.

The specification discloses that the claimed transgenic plants and methods can be used in breeding programs and expression assays. *See, e.g.*, Specification at page 18, lines 18-19, page 56, line 15 through page 75, line 10, and page 44, line 20 through page 48, line 21. The Examiner asserts “there is no art of record that discloses or provides any evidence that points to an activity for SEQ ID NO: 1 or its corresponding full length cDNA or the proteins that might be obtained using the full length cDNA to be obtained, such that another non-asserted utility would be well established.” *Id.*

The Examiner further argues that “Applicants have not identified any practical benefit of either the claimed plant or methods since neither the specification as filed, nor any prior art teaches or provides any evidence that defines any beneficial biological activity for the nucleic acid that is transformed into the plant, or assayed for in the claimed methods.” *Id.* at page 5.

This analysis misstates the nature of the asserted uses, ignores disclosed utilities, and misapplies the doctrine of “practical utility” developed by the courts after *Brenner v. Manson*. The “threshold for utility is not high: An invention is ‘useful’ under section 101 if it is capable of providing some identifiable benefit.” *Juicy Whip, Inc. v. Orange Bang, Inc.*, 185 F.3d 1364, 1366, 51 U.S.P.Q.2d 1700, 1702 (Fed. Cir. 1999), *citing Brenner v. Manson*, 383 U.S. 519, 534 (1966).

The Federal Circuit has recently reiterated that the “basic *quid pro quo* contemplated by the Constitution and the Congress for granting a patent monopoly is the benefit derived from the public from an invention with *substantial utility*.” *In re Fisher*, 421 F.3d 1365, -- U.S.P.Q.2d -- (Fed. Cir. 2005) (citing *Brenner*, 383 U.S. at 534-35) (emphasis in original). The Court noted that since “*Brenner* our predecessor court, the Court of Customs and Patent Appeals, and this court have required a claimed invention to have a specific and substantial utility to satisfy § 101” *Id.*

Although the Supreme Court has not defined the meaning of the terms “specific” and “substantial”, the Federal Circuit discerned the kind of disclosure an application could contain to establish a specific and substantial utility. *Id.* First, the Court indicated that the specification disclose a utility such that “one skilled in the art can use a claimed discovery in a manner which provides some *immediate benefit to the public*.” *Id.* (emphasis original). Second, the specification should also disclose “a use which is not so vague as to be meaningless,” that is that the claimed invention “can be used to provide a well-defined and particular benefit to the public.” *Id.*

It is well-established that claims must be considered as a whole in determining compliance with § 101. *Diamond v. Diehr*, 450 U.S. 175, 188, 209 U.S.P.Q. 1, 9 (1981). It is inappropriate to dissect claims and consider some elements while ignoring others. *Id.* Further, it is well-established law that “when a properly claimed invention meets at least one stated objective, utility under section 101 is clearly shown.” *Raytheon Co. v. Roper Corp.*, 724 F.2d 951, 958, 220 U.S.P.Q. 592, 598 (Fed. Cir. 1983).

Appellant has asserted in the specification that the claimed transgenic plants and methods provide clear and immediate benefits, for example, use to follow a plant through a

breeding program (*see, e.g.*, specification at page 18, lines 18-19, page 56, line 15 through page 75, line 10), and to determine the level or pattern of expression of a protein or mRNA associated with that nucleic acid molecule (*see, e.g.*, specification at page 44, line 20 through page 48, line 21). Either of these utilities described alone is enough to satisfy Section 101. Because Applicants need only establish a single utility to satisfy 35 U.S.C. § 101, and have done so in the present case, the premise of the rejection under Section 101 is incorrect, and the rejection should be reversed.

(1) The Claimed Plants and Methods Provide A Specific Benefit, *i.e.*, They Have Specific Utility

The specification describes multiple utilities for the present invention, including in breeding programs and in expression assays. Such uses provide a well-defined use that is not so vague as to be meaningless.

(a) Use of Transgenic Plants in Breeding Programs

For example, one of the utilities disclosed in the specification is use of the claimed transgenic plants in breeding programs (*see, e.g.*, specification at page 18, lines 18-19, page 56, line 15 through page 75, line 10). The Examiner rejects this asserted utility apparently because “neither the specification as filed, nor the prior art teaches or provides any evidence that defines any beneficial biological activity for the nucleic acid.” Final Action at page 5. Contrary to the Examiner’s position, the transformed plants having, *inter alia*, a structural nucleic acid molecule comprising a nucleic acid sequence of SEQ ID NO:1 or complement thereof, have utility independent of whether a function is known for the nucleic acid sequence. The specification discloses methods for the preparation of the transgenic plants as well as use in breeding programs to produce plants having genes of interest. *See, e.g.*, specification at page 18, lines 18-19 and page 56, line 15 through page 75, line 10. The specification further describes that the nucleic acid sequences can be used as markers. *See, e.g.*, specification at page 48, line 22 through page 49, line 5. The skilled artisan would recognize that such transformed plants can be more easily followed through a breeding program by the detection of the nucleic acid molecule. These utilities are immediately apparent for the claimed plants and methods without the need for further research.

The Examiner argues that this utility is not specific or substantial, apparently because this utility is “directed to basic research that involves studying the properties of the claimed transformed plant.” Final Action at page 4. The Examiner also argues that this utility is not “specific since the recited use can be generally applied to any plant transformed with any cDNA molecule or fragment thereof.” *Id.* The use of the claimed transformed plants having a nucleic acid molecule comprising, *inter alia*, a structural nucleic acid molecule comprising a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 and complement thereof in plant breeding programs is not a use directed to “basic research.” Rather, using the claimed transformed plants in a breeding program allows the breeder to readily track the transformed plant through the program by identifying progeny plants containing the nucleic acid molecule. Such a use is a well-defined utility. For example, progeny resulting from a cross between a claimed transformed plant having a desired background and a plant having a trait of interest to be introgressed into the transformed plant can be screened quickly for identification of the nucleic acid molecule. Such a plant allows for the high-throughput screening of progeny plants to obtain the desired commercial plant.

The asserted use of the claimed plants in breeding programs is not a “nebulous” or “obscure” expression of utility. The claimed transformed plants have been asserted to work for a specific, *i.e.*, not vague or unknown benefit, for use in breeding programs. This benefit is immediately realized directly from the use of the claimed transformed plants, not from the use of other plants. Such a proven use that provides an acknowledged known benefit to the public satisfies the utility requirement of 35 U.S.C. § 101.

(b) Methods for Determining a Level or Pattern in a Plant Cell or Plant Tissue of a Protein in a Plant

Uses for the claimed methods include detecting the presence or absence or level of expression of the sequence in a sample (*See e.g.*, specification at page 44, line 20 through page 48, line 21). The Examiner suggests that these uses are not legal utilities because “further experimentation would be required to identify the full length cDNA comprising SEQ ID NO: 1, isolate the protein encoded by the full length cDNA, and furthermore identify the biological activity of the encoded protein.” Final Action, at page 6. This is not

correct. Such methods can be used, for example, to assay gene expression in plant cells treated with an herbicide to detect target genes for producing herbicide tolerant plants. The Examiner has not provided any evidence that would reasonably suggest that this cannot be done, and thus has not met the burden of proof required to establish a utility rejection. *See In re Brana*, 51 F.3d 1560, 1567, 34 U.S.P.Q.2d 1436, 1441 (Fed. Cir. 1995). *Accord In re Gaubert*, 524 F.2d 1222, 1225-26, 187 U.S.P.Q. 664, 666 (C.C.P.A. 1975); *In re Langer*, 503 F.2d 1380, 1391, 183 U.S.P.Q. 288, 297 (C.C.P.A. 1974).

One illustrative example of a molecule that can be detected using a claimed method is the nucleic acid sequence of SEQ ID NO: 1. Appellant has specifically disclosed that one use of the claimed method is to detect the level in a plant cell or plant tissue of a protein in a plant. Specification at page 44, line 20 through page 48, line 21. The Examiner denigrates that utility by asserting that it is not specific because it is generally applicable to any nucleic acid. Final Action at pages 4-5. This is not correct. The claimed methods using the nucleic acid molecules are particularly useful, for example, to detect the level in a plant cell or tissue of an mRNA corresponding to SEQ ID NO: 1. *See, e.g.*, specification at page 44, line 20 through page 48, line 21.

In short, the Examiner suggests that the asserted utilities are legally insufficient simply because other molecules can be used for the same purpose, *e.g.*, monitoring gene expression or in a breeding program. That position is wrong as a matter of law – there is no requirement of exclusive utility in the patent law. *See Carl Zeiss Stiftung v. Renishaw PLC*, 945 F.2d 1173, 1180, 20 U.S.P.Q.2d 1094, 1100 (Fed. Cir. 1991) (“An invention need not be the best or the only way to accomplish a certain result...”). Such an argument would imply that a new golf club has no legal utility because other golf clubs can be used for the same purpose, *i.e.*, hitting golf balls. That position must be rejected as it requires reading “into the patent laws limitations and conditions which the legislature has not expressed,” a practice condemned by the Supreme Court. *See Diamond v. Chakrabarty*, 447 U.S. 303, 308, 206 U.S.P.Q. 193, 196 (1980), *quoting United States v. Dubilier Condenser Corp.*, 289 U.S. 178, 199, 17 U.S.P.Q. 154, 162 (1933).

Moreover, it is factually incorrect that this use is not “specific” to the claimed plants and methods. The claimed methods and plants provide a particularly appropriate and demonstrably useful starting point for example, to screen for compounds with herbicidal activity. A random nucleic acid molecule does not provide an equally good starting point for such an assay as a random nucleic acid molecule would not necessarily correlate to an expressed gene. Furthermore, even if a random nucleic acid molecule provided a better starting point than the claimed nucleic acid molecules, it would not obviate the utility of the claimed plants and methods. An invention may be “less effective than existing devices but nevertheless meet the statutory criteria for patentability.” *Custom Accessories, Inc. v. Jeffrey-Allan Indus.*, 807 F.2d 955, 960 n.12, 1 U.S.P.Q.2d 1196, 1199 n.12 (Fed. Cir. 1986).

(c) Methods for Determining a Level or Pattern in a Plant Cell or Tissue of a Protein in a Plant using *In Situ* Hybridization

The specification discloses additional specific and substantial uses of the methods, for example for use in *in situ* hybridizations (*See, e.g.*, Specification at page 45, line 10 through page 47, line 11 and Claim 9). *In situ* hybridization is a highly sensitive technique that can detect as few as 5-10 copies of an RNA sequence per cell and generally involves tissue preparation, hybridization, and washing conditions, and can be used, *inter alia*, to map chromosomal locations of genes, and as diagnostic tools. *See, e.g.*, Specification at page 45, line 10 through page 47, line 11. *In situ* hybridization can be used in a number of uses, for example to determine the spatial population or the steady-state levels of RNA accumulation in a tissue. One of the most common uses of *in situ* hybridization is to localize specific RNA sequences in cells, which is useful for gene mapping, following chromosomes in hybrid lines or detecting chromosomes with translocations, transversions, or deletions. As such, methods for determining the level or pattern in a plant cell or plant tissue of a protein using *in situ* hybridization provides a useful tool for studying cellular processes.

Many of the disclosed utilities in this case, including *in situ* hybridization, are directly analogous to the utilities of a microscope, *i.e.*, the claimed methods employing the recited nucleic acid molecules may be used to locate and measure nucleic acid molecules

within a sample, cell, or organism. The Examiner denigrates such utilities by asserting that these utilities are not “useful” because “[w]ithout identifying the biological activity of the encoded protein, the skilled artisan would not be able to recognize the real-world context of use of the” claimed methods. Final Action, at page 6. However, the fact that, *e.g.*, a new and nonobvious microscope or screening assay can be used for learning about products or processes does not lessen the fact that such “tools” have legal utility. “Many research tools such as gas chromatographs, screening assays, and nucleotide sequencing techniques have clear, specific and unquestionable utility (*e.g.*, they are useful in analyzing compounds).” MPEP § 2107.01 at page 2100-33.

Use of the claimed methods to determine the level or pattern of a protein is no more legally insufficient than using a gas chromatograph to analyze the chemical composition of a gas – such use determines information about the gas, not the gas chromatograph. Even if the gas chromatograph detects the absence of a particular chemical element in the gas, that finding does not obviate the utility of the gas chromatograph itself. Information has been obtained about the gas.⁴ Likewise, the claimed methods have utility even if the absence of a particular molecule is detected. Indeed, the absence of the corresponding molecule usefully demonstrates that the nucleic acid molecule is not expressed in a given cell or tissue.

(d) Methods for Determining a Level or Pattern in a Plant Cell or Tissue of a Protein in a Plant using Tissue Printing

The specification also discloses that the methods can be used to detect the level or pattern of a protein using tissue printing (*See*, Claim 10). Tissue printing provides a convenient method to simultaneously screen on a single membrane many tissue sections from different plants or different developmental stages. Tissue printing procedures utilize various films designed to immobilize proteins and nucleic acids. Briefly, a freshly cut section of a plant tissue or organ is pressed gently onto nitrocellulose paper, nylon membrane, or

⁴ For example, gas sampled from crude oil may be analyzed by gas chromatography for the presence or absence of chlorine, which is toxic to catalysts used in gasoline refining even in very low concentrations. The absence of a peak at the molecular weight of chlorine indicates the absence of chlorine in the sample being tested, thereby providing useful information (no chlorine is present, therefore the catalyst will not be destroyed) to the refinery manager. *See, e.g.*, U.S. Patent No. 6,133,740 entitled “Chlorine Specific Gas Chromatographic Detector.”

polyvinylidene difluoride membrane. Such methods can be used to visualize RNA accumulation in tissues and organs and during different developmental stages. The specification discloses that tissue printing can be used for the histochemical localization of various plant enzymes and nucleic acids. *See, e.g.* Specification at page 47, line 12 through page 48, line 17.

Such utility, as are many of the disclosed utilities, is also directly analogous to the utilities of a microscope, *i.e.*, the claimed methods employing the recited nucleic acid molecules in tissue printing may be used to spatially and temporally localize proteins and nucleic acid molecules within a sample, cell, or organism.

The Examiner has failed to provide evidence, or even to suggest a reason for believing that the claimed plants and methods could not be so used. Accordingly, the assertion of this utility in a breeding program or in expression assays satisfies the requirements of 35 U.S.C. § 101. *See In re Brana*, 51 F.3d 1560, 1566, 34 U.S.P.Q.2d 1436, 1441 (Fed. Cir. 1995).

(2) The Claimed Transgenic Plants and Methods Provide Practical, Real World Benefits, *i.e.*, They Have Substantial Utility

The Final Action also appears to assert that the disclosed uses for the claimed plants or the claimed methods are legally insufficient because they are not “substantial” utilities. Final Action at pages 2-6. The touchstone of “substantial” utility is “real world” or “practical utility.” *See, e.g., Fujikawa v. Wattanasin*, 93 F.3d 1559, 1563, 39 U.S.P.Q.2d 1895, 1899 (Fed. Cir. 1996). “ ‘Practical utility’ is a shorthand way of attributing ‘real world’ value to claimed subject matter. In other words, one skilled in the art can use a claimed discovery in a manner which provides some immediate benefit to the public.” *Nelson v. Bowler*, 626 F.2d 853, 856, 857, 206 U.S.P.Q. 881, 883 (C.C.P.A. 1980) (“tests evidencing pharmacological activity may manifest a practical utility even though they may not establish a specific therapeutic use”).⁵

⁵ *Accord Cross v. Iizuka*, 753 F.2d 1040, 1050, 224 U.S.P.Q. 739, 747-48 (Fed. Cir. 1985); *Rey-Bellet v. Engelhardt*, 493 F.2d 1380, 1383, 181 U.S.P.Q. 453, 454 (C.C.P.A. 1974).

There can be no question that one skilled in the art can use the claimed transgenic plants and methods in a manner which provides an immediate benefit to the public, for example to easily identify progeny of interest in a breeding program. The detection of transgenic plants provides an immediate benefit to the public because, *e.g.*, it enables a plant breeder to more efficiently allocate resources to progeny plants having a given genetic profile. This information about a plant's genetic profile, like the information about a compound's pharmacological profile in *Nelson*, provides an immediate benefit and thus a practical utility to the public.

(3) The Disclosed Utilities Are Credible to One of Skill in the Art

An assertion of utility must be accepted by the Examiner unless it would not be considered "credible" by a person of ordinary skill in the art. MPEP § 2107 at 2100-29. Cases in which utility was found not to be credible are rare, and usually involve "hare-brained" utilities.⁶ A challenge to the credibility of a utility is essentially a challenge directed to operability, and such a challenge must be supported by a clear statement of "factual reasons which would lead one skilled in the art to question the objective truth of the statement of operability." *In re Gaubert*, 524 F.2d 1222, 1225-26, 187 U.S.P.Q. 664, 666 (C.C.P.A. 1975); *see In re Brana*, 51 F.3d 1560, 1567, 34 U.S.P.Q.2d 1436, 1441 (Fed. Cir. 1995); MPEP § 2107.02 at 2100-41.

Applicants have explicitly identified specific and substantial utilities, not only in the specification, but in Applicants' responses. "To violate [35 U.S.C.] 101 the claimed

⁶ Examples of incredible utilities are given in MPEP § 2107.01 at page 2100-34, and include:

an invention asserted to change the taste of food using a magnetic field (*Fregeau v. Mossinghoff*, 776 F.2d 1034, 227 U.S.P.Q. 848 (Fed. Cir. 1985)), a perpetual motion machine (*Newman v. Quigg*, 877 F.2d 1575, 11 U.S.P.Q. 1340 (Fed. Cir. 1989)), a flying machine operating on "flapping or flutter function" (*In re Houghton*, 433 F.2d 820, 167 U.S.P.Q. 687 (C.C.P.A. 1970)), a method for increasing the energy output of fossil fuels upon combustion through exposure to a magnetic field (*In re Ruskin*, 354 F.2d 395, 148 U.S.P.Q. 221 (C.C.P.A. 1966)), uncharacterized compositions for curing a wide array of cancers (*In re Citron*, 325 F.2d 248, 139 U.S.P.Q. 516 (C.C.P.A. 1963)), a method of controlling the aging process (*In re Eltgroth*, 419 F.2d 918, 164 U.S.P.Q. 221 (C.C.P.A. 1970)), and a method of restoring hair growth (*In re Ferens*, 417 F.2d 1072, 163 U.S.P.Q. 609 (C.C.P.A. 1969)).

device must be totally incapable of achieving a useful result.” *Brooktree Corp. v. Advanced Micro Devices, Inc.*, 977 F.2d 1555, 1571, 24 U.S.P.Q.2d 1401, 1412 (Fed. Cir. 1992). To date, the Examiner has provided no evidence that the claimed plants and methods will not work for the asserted utilities. Unless and until the Examiner can prove that the claimed invention is wholly inoperative, the rejection must be withdrawn.

In view of the above, Applicants contend that the claimed plants and methods are supported by credible, specific, and substantial utilities disclosed in the specification. Moreover, the Examiner has failed to raise any credible evidence challenging the presently asserted utilities. Consequently, the rejection of pending claims 4-12 under 35 U.S.C. §101 is improper and should be reversed.

C. The Claimed Plants and Methods Are Enabled by the Specification

The enablement of the claimed transgenic plants and methods has been challenged. Claims 4-12 have been erroneously rejected as not enabled by the specification, because the claimed nucleic acid molecules allegedly lack utility and therefore cannot be enabled. Final Action at page 7. This rejection is erroneous and has been overcome by the arguments stated above regarding utility because it is well-established law that “the enablement requirement is met if the description enables any mode of making and using the invention.” *Johns Hopkins University v. CellPro*, 152 F.3d 1342, 1361, 47 U.S.P.Q.2d 1705, 1719 (Fed. Cir. 1998) (emphasis added), *quoting Engel Indus. v. Lockformer Co.*, 946 F.2d 1528, 1533, 20 U.S.P.Q.2d 1300, 1304 (Fed. Cir. 1991). Unless and until the Examiner comes forth with evidence to rebut the objective truth of the utilities disclosed in the specification, this enablement rejection must be withdrawn as improper. *See In re Wright*, 999 F.2d 1557, 1561-62, 27 U.S.P.Q.2d 1510, 1513 (Fed. Cir. 1993); *Ex parte Lemak*, 210 U.S.P.Q. 306, 307 (Bd. App. 1981) (“pure conjecture” does not substantiate rejection for lack of enablement).

D. The Specification Provides An Adequate Written Description of the Claimed Invention

The adequacy of the written description of the claimed invention of claims 4-12 has been challenged by the Examiner because the claimed subject matter was allegedly “not

described in the specification in such a way as to reasonably convey to one skilled in the relevant art that the inventor(s)...had possession of the claimed invention.” Final Action at page 7. The Examiner contends that the rejection has been maintained “for the reasons of record set forth in the Examiner’s Answer mailed 10/23/2002.” Final Action at page 7. The Examiner rejects the claims on the basis of “the breadth of the claims are directed to plants transformed with nucleic acids encompassing full length gene sequences (i.e. gene sequences yet to be discovered) and cDNAs comprising SEQ ID NO: 1, sequences that hybridize to SEQ ID NO: 1, and methods which utilize such sequences” Final Action at page 8. However, the specification demonstrates to one skilled in the art that Applicants were in possession of the claimed invention containing the genera of nucleic acid molecules.

(1) The Specification Reflects Applicants’ Possession of the Claimed Invention

The purpose of the written description requirement is to ensure that the inventor had possession of the claimed subject matter, *i.e.*, to ensure that the inventors actually invented what is claimed. *Gentry Gallery Inc. v. Berkline Corp.*, 134 F.3d 1473, 1479, 45 U.S.P.Q.2d 1498, 1503 (Fed. Cir. 1998); *Lockwood v. American Airlines*, 107 F.3d 1565, 1572, 41 U.S.P.Q.2d 1961, 1966 (Fed. Cir. 1997); *In re Alton*, 76 F.3d 1168, 1172, 37 U.S.P.Q.2d 1578, 1581 (Fed. Cir. 1996). If a person of ordinary skill in the art would, after reading the specification, understand that the inventors had possession of the claimed invention, even if not every nuance, then the written description has been met. *In re Alton*, 76 F.3d at 1175, 37 U.S.P.Q.2d at 1584. A person of ordinary skill in the art would, after reading the present specification, understand that Applicants had possession of transgenic plants and methods employing nucleic acid molecules comprising or specifically hybridizing to a nucleic acid sequence of SEQ ID NO: 1. Applicants have provided the nucleotide sequence required by the claims, *e.g.*, SEQ ID NO: 1. Accordingly, Applicants have demonstrated possession of the claimed invention.

The fact that the claims at issue are intended to include molecules that encode the recited enzymes or fragments of the recited nucleic acid sequence, the recited sequence joined with additional sequences, related sequences or complements of the recited

sequence, does not mean that Applicants were any less in possession of the nucleic acid molecules.⁷ It is well-established law that use of the transitional term “comprising” properly leaves the claims “open for the inclusion of unspecified ingredients even in major amounts.” *Ex parte Davis*, 80 U.S.P.Q. 448, 450 (BPAI 1948). *Accord PPG Indus. v. Guardian Indus.*, 156 F.3d 1351, 1354, 48 U.S.P.Q.2d 1351, 1353-54 (Fed. Cir. 1998); *Moleculon Research Corp. v. CBS*, 793 F.2d 1261, 1271, 229 U.S.P.Q. 805, 812 (Fed. Cir. 1986).

The present application describes more than just the nucleotide sequence recited by the claims (*e.g.* SEQ ID NO: 1). For example, the specification describes the transformation of plants with vectors having the claimed nucleic acid molecules (*See, e.g.*, specification at page 66, line 12 through page 75, line 10) as well as methods for determining gene expression (*See, e.g.*, specification at page 15, lines 3-16 and page 47, line 8 through page 50, line 15). Moreover, the present application describes more than just the nucleotide sequence used in the claims (SEQ ID NO: 1). For example, it describes vectors comprising the nucleic acid molecules, Specification at page 56, line 15 through page 66, line 11, and describes how to make the nucleotide sequences and the libraries from which they were originally purified. *See* specification at page 1, line 16 through page 4, line 23, and Examples 1-2.

Moreover, the court determined, in *Enzo Biochem, Inc. v. Gen-Probe Inc.*, 296 F.3d 1316, 1321, 63 U.S.P.Q.2d 1609, 1610 (Fed. Cir. 2002), that the written description inquiry is a factual one determined on a case-by-case basis and that, in a given disclosure, “it may well be that various subsequences, mutations, and mixtures of those sequences are also described to one of skill in the art.” *Enzo*, 296 F.3d at 1326-1327, 63 U.S.P.Q.2d at 1615. Furthermore, it is well established that claims “may be broader than the specific embodiment disclosed in a specification.” *Ralston-Purina Co. v. Far-mor-Co*, 772 F.2d

⁷ If the Examiner is arguing that no possession is shown because the precise claim language is not used in the specification, then it goes beyond what is required by the law. It is well-settled that the description of a claimed invention need not be *in ipso verbis*. *Gentry Gallery v. Berkline Corp.*, 134 F.3d 1473, 1479, 45 U.S.P.Q.2d 1498, 1503 (Fed. Cir. 1998); *In re Alton*, 76 F.3d 1168, 1175, 37 U.S.P.Q.2d 1578, 1583 (Fed. Cir. 1996); *Martin v. Johnson*, 454 F.2d 746, 751, 172 U.S.P.Q. 391, 395 (C.C.P.A. 1972).

1570, 1575, 227 U.S.P.Q. 177, 179 (Fed. Cir. 1985) (*quoting In re Rasmussen*, 650 F.2d 1212, 1215, 211 U.S.P.Q. 323, 326 (C.C.P.A. 1981)). A complete structure of every species within a chemical genus is not required. *See, e.g., Utter v. Hiraga*, 845 F.2d 993, 998, 6 USPQ2d 1709, 1714 (Fed. Cir. 1988) (“A specification may, within the meaning of 35 U.S.C. § 112, ¶ 1, contain a written description of a broadly claimed invention without describing all species that claim encompasses.”).

The Examiner asserts that the claimed invention covers “gene sequences yet to be discovered”, and accordingly Appellant has allegedly not adequately disclosed the genera of nucleic acid molecules. Final Action at page 8. As such, the Examiner appears to require that each nucleic acid molecule within the genera must be described by its complete structure. This requirement is totally unfounded. The Federal Circuit has elucidated a test for written description wherein a genus of nucleic acids may be described by a structural feature that distinguishes members of the claimed genus from non-members of the claimed genus. *Regents of the University of California v. Eli Lilly and Co.*, 119 F.3d 1559, 1568-69, 43 U.S.P.Q.2d 1398, 1406 (Fed. Cir. 1997). Applicants have satisfied that test for written description.

In particular, Applicants have disclosed common structural features of the nucleic acid molecule recited in claim 4, for example the nucleotide sequence of SEQ ID NO: 1. The respective common structural feature (the nucleotide sequence of SEQ ID NO: 1) is shared by every nucleic acid molecule in the claimed genera, and it distinguishes the members of the claimed genera from non-members. For example, if a nucleic acid molecule such as an mRNA contains the nucleotide sequence of SEQ ID NO: 1, then it is a member of the genus of nucleic acid molecules comprising a nucleic acid sequence of SEQ ID NO: 1. If a nucleic acid molecule does not contain SEQ ID NO: 1, then it is not a member of that claimed genus. The presence of other nucleotides at either end of the recited sequence will not interfere with the recognition of a claimed nucleic acid molecule as such – it either contains the nucleotides of SEQ ID NO: 1 or it does not. One skilled in the art, after reading the present specification, would clearly know if a nucleic acid molecule contains one of the recited nucleotide sequences. Similarly, if a nucleic acid

molecule specifically hybridizes to the nucleic acid sequence of SEQ ID NO: 1, then it is a member of the genus of nucleic acid molecules recited in claim 8. If the nucleic acid molecule does not specifically hybridize, then it is not a member of that genus.

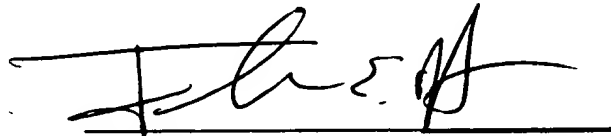
Thus, claims 4-12 are supported by an adequate written description pursuant to the requirements of 35 U.S.C. § 112, and the rejection should be reversed.

CONCLUSION

In view of the foregoing, it is respectfully requested that the Board of Patent Appeals and Interferences reverse the Rejections and that the subject application be allowed forthwith.

Respectfully submitted,

Date: September 25, 2006



Thomas E. Holsten (Reg. No. 46,098)
David R. Marsh (Reg. No. 41,408)

Of Counsel
Lawrence M. Lavin, Jr. (Reg. No. 30,768)
Thomas E. Kelley (Reg. No. 29,938)
Monsanto Company

ARNOLD & PORTER LLP
Attn: IP Docketing
555 Twelfth Street, NW
Washington, DC 20004-1206
202.942.5000 telephone
202.942.5999 facsimile

CLAIMS APPENDIX

Claims 1-3 (Cancelled)

4. (Previously presented) A transformed plant having a nucleic acid molecule which comprises:
 - (a) an exogenous promoter region which functions in a plant cell to cause the production of an mRNA molecule;
 - (b) a structural nucleic acid molecule comprising a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 and complement thereof; and
 - (c) a 3' non-translated sequence that functions in said plant cell to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of said mRNA molecule.
5. (Previously presented) The transformed plant according to claim 4, wherein said structural nucleic acid molecule is a complement of the nucleic acid sequence of SEQ ID NO: 1.
6. (Original) The transformed plant according to claim 5, wherein said plant is soybean or maize.
7. (Original) The transformed plant according to claim 5, wherein said plant is soybean.
8. (Previously presented) A method for determining a level or pattern in a plant cell or plant tissue of a protein in a plant comprising:

- (a) incubating, under conditions permitting nucleic acid hybridization, a marker nucleic acid molecule, said marker nucleic acid molecule selected from the group of marker nucleic acid molecules which specifically hybridize to a nucleic acid molecule having the nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 or complement thereof, with a complementary nucleic acid molecule obtained from said plant cell or plant tissue, wherein nucleic acid hybridization between said marker nucleic acid molecule and said complementary nucleic acid molecule obtained from said plant cell or plant tissue permits the detection of an mRNA for said protein;
 - (b) permitting hybridization between said marker nucleic acid molecule and said complementary nucleic acid molecule obtained from said plant cell or plant tissue; and
 - (c) detecting the level or pattern of said complementary nucleic acid, wherein the detection of said complementary nucleic acid is predictive of the level or pattern of said protein.
9. (Original) The method of claim 8, wherein said level or pattern is detected by *in situ* hybridization.
10. (Original) The method of claim 8, wherein said level or pattern is detected by tissue printing.
11. (Original) The method of claim 8, wherein said plant is maize or soybean.
12. (Original) The method of claim 11, wherein said plant is soybean.

Claims 13-17 (Cancelled)

EVIDENCE APPENDIX

No New Evidence

RELATED PROCEEDINGS APPENDIX

Attached are copies of Decisions on Appeal that issued in the following applications:

1. *In re Fisher*, 412 F.3d 1365 (Fed. Cir. 2005)
2. U.S. Appln. No. 09/619,643, BPAI Appeal No. 2002-2046
3. U.S. Appln. No. 09/654,617, BPAI Appeal No. 2003-1744
4. U.S. Appln. No. 09/620,392, BPAI Appeal No. 2003-1746
5. U.S. Appln. No. 09/540,232, BPAI Appeal No. 2003-1137
6. U.S. Appln. No. 09/440,687, BPAI Appeal No. 2003-1504
7. U.S. Appln. No. 09/565,240, BPAI Appeal No. 2003-1135
8. U.S. Appln. No. 09/540,215, BPAI Appeal No. 2003-0996
9. U.S. Appln. No. 09/552,087, BPAI Appeal No. 2004-1772
10. U.S. Appln. No. 09/206,040, BPAI Appeal No. 2002-0078

Ex parte Fisher

U.S. Patent and Trademark Office Board of Patent
Appeals and Interferences

Appeal No. 2002-2046

Decided April 1, 2004

PATENTS

[1] Patentability/Validity -- Utility (§ 115.10)

Invention must have "substantial" utility in order to be "useful" within meaning of 35 U.S.C. § 101, since, if Section 101's requirement is given its broadest reach, then little or nothing of chemical nature would be found to lack utility; thus, invention must provide specific benefit in its currently available form, and vague, general disclosures or arguments that invention is "useful in research," or statements that invention is capable of being used for particular purpose, would not satisfy Section 101.

[2] Patentability/Validity -- Utility (§ 115.10)

Disclosed use of claimed nucleic acid molecules encoding maize protein in identifying and detecting polymorphisms in population of maize plants does not provide "substantial" degree of utility required for invention to be "useful" within meaning of 35 U.S.C. § 101, since specification does not explain why claimed nucleotide molecules would in fact be useful in detecting polymorphisms, and since, without further information regarding gene represented by molecules, detection of presence or absence of polymorphism provides only barest information concerning genetic heritage.

[3] Patentability/Validity -- Utility (§ 115.10)

Disclosed use of claimed nucleic acid molecules encoding maize protein to isolate nucleic acid molecules of other plants and organisms does not provide "substantial" degree of utility required for invention to be "useful" within meaning of 35 U.S.C. § 101, since specification does not attribute any property, in terms of plant trait or phenotype, to any of disclosed nucleotide molecules, and absent such information, using claimed molecules to isolate other molecules does not represent substantial utility; applicants' assertion that claimed nucleic acid molecules may be useful in searching for certain promoters does not supply required utility, since specification does not provide any expectation of successfully using any of claimed molecules to isolate such promoters, and since there is no evidence that claimed molecules are tissue or cell-type specific, or developmentally or environmentally regulated.

[4] Patentability/Validity -- Utility (§ 115.10)

Invention can have utility shared by other compounds or compositions, but not every utility will satisfy requirements of 35 U.S.C. § 101, even if utility is shared by class of inventions; thus, although utility need not be unique to claimed invention, it must nevertheless be specific, and in currently available form, in order to satisfy Section 101.

[5] Patentability/Validity -- Utility (§ 115.10)

Assertion that claimed nucleic acid molecules encoding maize protein can be introduced into plant or plant cell, which can then be used to screen for compounds such as herbicides, does not provide "substantial" degree of utility required for invention to be "useful" within meaning of 35 U.S.C. § 101, since invention claims otherwise uncharacterized nucleic acid molecule, not "cell-based assay," and since specification does not indicate that asserted screening method is feasible if uncharacterized nucleic acid is used.

[6] Patentability/Validity -- Utility (§ 115.10)

Disclosed use of claimed nucleic acid molecules encoding maize protein as components of assays for monitoring gene expression does not provide "substantial" degree of utility required for invention to be "useful" within meaning of 35 U.S.C. § 101, since specification provides no guidance that would allow person skilled in art to use data relating to expression of gene in any practical way; claim that those skilled in art could experiment with claimed nucleic acids and figure out for themselves what results might mean does not provide "specific benefit in currently available form," and although each nucleic acid used in gene expression assay contributes to data generated by assay as whole, fact that assay itself may have patentable utility does not mean that each tiny component of assay also has patentable utility.

[7] Patentability/Validity -- Specification -- Written description (§ 115.1103)

Application claiming five nucleic acid molecules encoding maize protein was improperly rejected for failure to satisfy written description requirement of 35 U.S.C. § 112, since application provides nucleotide sequences for five molecules, and claim's use of "comprising" allows for addition of nucleotides or other molecules at either end of nucleotide sequences without permitting internal alterations of sequences, and since fact that claimed molecules may have other molecules attached at their ends does not diminish adequate written description of molecules having sequences set forth in application.

*1021 Patent application of Dane K. Fisher and Raghunath V. Lalgudi, serial no. 09/619,643 (nucleic

acid molecule encoding maize protein or fragment thereof). Applicants appeal from examiner's final rejection of sole claim pending in application. Affirmed in part and reversed in part.

[Editor's Note: The Board of Patent Appeals and Interferences states that this opinion was not written for publication, and is not binding precedent of the board.]

Lawrence M. Lavin Jr., Monsanto Co., St. Louis, Mo., for applicants.

Before William F. Smith, Adams, and Grimes, administrative patent judges.

Adams, J.

DECISION ON APPEAL

This is a decision on the appeal under 35 U.S.C. § 134 from the examiner's final rejection of claim 1, the only claim pending in the application, reproduced below:

1.A substantially purified nucleic acid molecule that encodes a maize protein or fragment thereof comprising a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO:5.

The examiner does not rely on a reference.

GROUND OF REJECTION

Claim 1 stands rejected under 35 U.S.C. § 101 as lacking utility and § 112, first paragraph, for lack of enablement based on the finding of lack of utility. Claim 1 also stands rejected under 35 U.S.C. § 112, first paragraph, as the specification fails to provide an adequate written description of the claimed invention. We affirm the utility and enablement rejections. We reverse the written description rejection.

BACKGROUND

The subject matter of the present appeal is directed to expressed sequence tags (ESTs). *See* Specification, page 15, lines 9-10. ESTs "are short sequences of randomly selected clones from a cDNA (or complementary DNA) library which are representative of the cDNA inserts of these randomly selected clones." Specification, page 1.

As set forth at page 9, lines 2-4, of appellants' specification "[t]he present invention provides a substantially purified nucleic acid molecule that encodes a maize protein or fragment thereof comprising a nucleic acid sequence selected from the

group consisting of *1022 SEQ ID NO: 1 through SEQ ID NO: 32236." Of these 32,236 nucleic acid sequences, the originally filed claims were directed to SEQ ID NO: 1 through SEQ ID NO: 4,013. On January 26, 2001 (Paper No. 4), the examiner entered a Restriction requirement into the record, requiring, *inter alia*, appellants "to elect up to 5 nucleic acid sequences" for consideration on the merits. Paper No. 4, page 3. In response, appellants elected SEQ ID NO:1 through SEQ ID NO:5. The ESTs set forth in SEQ ID NO: 1 through SEQ ID NO: 5 are disclosed to be obtained from cDNA library LIB3115 "generated from maize (RX601, Asgrow Seed Company, Des Moines, Iowa U.S.A.) pooled leaf tissue. . . ." Specification, pages 79-80, Example 1.

The specification sets forth a number of utilities for the nucleic acid molecules of SEQ ID NO: 1 through SEQ ID NO: 5 which are summarized by the examiner (Answer, bridging paragraph, pages 5-6) as follows:

The specification teaches that the nucleic acids may be used to produce a plant containing reduced levels of a protein (pg. 11), determining an association between a polymorphism and a plant trait (pg. 11), isolating a genetic region or nucleic acid (pg. 11), determining a level or pattern in a plant cell of a protein in a plant (pg. 11), determining a mutation in a plant whose presence is predictive of a mutation affecting a level or pattern of a protein (pg. 13), as molecular tags to isolate genetic regions, isolate genes, map genes, and determine gene function (pg. 14), and identifying tissues (pg. 14)[.] The specification states that the nucleic acid ESTs of the present invention can enable the acquisition of molecular markers, which can be used in breeding schemes, genetic and molecular mapping and cloning of agronomically significant genes (pg. 31).

In the examiner's opinion "[t]hese are non-specific uses that are applicable to nucleic acids in general and not particular or specific to the nucleic acids being claimed." Answer, page 6. For example, the examiner finds (Answer, page 10), "determining whether the claimed nucleic acids have or do not have a polymorphism would require determining whether there was a polymorphism within such a sequence and then determining how to use this information in a patentably meaningful way." [FN1]

In presenting their case on appeal, appellants focus on use of the claimed nucleic acid molecules to identify the presence or absence of a polymorphism, and their use as probes or as a source for primers. *See e.g.*, Brief, pages 6- 12. According to appellants

(Brief, page 3), "they have disclosed nucleic acid molecules which, in their current form, provide at least one specific benefit to the public, for example the ability to identify the presence or absence of a polymorphism in a population of maize plants." Furthermore, appellants assert (Brief, page 8), "[t]he specification discloses that the claimed nucleic acid molecules can be used to isolate nucleic acid molecules of other plants and organisms. . . ."

CLAIM CONSTRUCTION

As set forth above, claim 1 on appeal is drawn to a substantially purified nucleic acid molecule that encodes a maize protein or fragment thereof comprising a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO:5. According to appellants' specification (page 15, lines 19-25), the term "substantially purified"

refers to a molecule separated from substantially all other molecules normally associated with it in its native state. More preferably a substantially purified molecule is the predominant species present in a preparation. A substantially purified molecule may be greater than 60% free, preferably 75% free, more preferably 90% free, and most preferably 95% free from the other molecules (exclusive of solvent) present in the natural mixture. The term "substantially purified" is not intended to encompass molecules present in their native state.

As we understand the claimed invention the use of the transitional term "comprising" does not allow for internal alterations (e.g. insertions or deletions) of the nucleotide sequences set forth in SEQ ID NO: 1 through SEQ ID NO: 5, but instead only allows for the addition of nucleotides or other molecules at either end of the nucleotide sequences set forth in SEQ *1023 ID NO: 1 through SEQ ID NO: 5. [FN2] In this regard, we recognize, as does the examiner (Answer, page 14), the claim as written encompasses, *inter alia*, genes, full open reading frames, fusion constructs, and cDNAs.

Accordingly, for the purposes of our review, we interpret the claimed invention as drawn to a nucleic acid molecule, separated from substantially all other molecules normally associated with it in its native state, selected from the group consisting of the nucleic acid molecule defined by the 429 nucleotide sequence set forth in SEQ ID NO: 1, the 413 nucleotide sequence set forth in SEQ ID NO: 2, the 365 nucleotide sequence set forth in SEQ ID NO: 3, the 414 nucleotide sequence set forth in SEQ ID NO: 4, and the 333 nucleotide sequence set forth in SEQ ID NO: 5, with or without any preceding or trailing

nucleotides, or other molecules.

DISCUSSION

Utility

The starting point for determining whether a nucleic acid molecule selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 5 possesses utility under 35 U.S.C. § 101 is Brenner v. Manson, 383 U.S. 519, 148 USPQ 689 (1966). As set forth in Brenner, at 534-35, 148 USPQ at 695 [FN3] ,

the basic quid pro quo contemplated by the Constitution and the Congress for granting a patent monopoly is the benefit derived by the public from an invention with substantial utility. Unless and until [an invention] is refined and developed to this point--where specific benefit exists in currently available form--there is insufficient justification for permitting an applicant to engross what may prove to be a broad field.

In considering the issues presented in this appeal, special attention must be paid to the Brenner court's statement that a patent should issue only when an invention possesses "substantial utility," *i.e.*, "where a specific benefit exists in currently available form." Whether a claimed invention is useful under 35 U.S.C. § 101 is a question of fact. Cross v. Iizuka, 753 F.2d 1040, 1044 n.7, 224 USPQ 739, 742 n.7 (Fed. Cir. 1985).

At issue in Brenner was a claim to "a chemical process which yields an already known product whose utility--other than as a possible object of scientific inquiry--ha[d] not yet been evidenced." Id. at 529, 148 USPQ at 693. The Patent Office had rejected the claimed process for lack of utility, on the basis that the product produced by the claimed process had not been shown to be useful. See id. at 521-22, 148 USPQ at 690. On appeal, the Court of Customs and Patent Appeals reversed, on the basis that "where a claimed process produces a known product it is not necessary to show utility for the product." Id. at 522, 148 USPQ at 691.

The Brenner Court noted that although § 101 requires that an invention be "useful," that "simple, everyday word can be pregnant with ambiguity when applied to the facts of life." Id. at 529, 148 USPQ at 693. Thus,

[i]t is not remarkable that differences arise as to how the test of usefulness is to be applied to chemical processes. Even if we knew precisely what Congress meant in 1790 when it devised the "new and useful"

phraseology and in subsequent re-enactments of the test, we should have difficulty in applying it in the context of contemporary chemistry, where research is as comprehensive as man's grasp and where little or nothing is wholly beyond the pale of "utility"--if that word is given its broadest reach.

Id. at 530, 148 USPQ at 694. [FN4]

The Court, finding "no specific assistance in the legislative materials underlying § 101," based its analysis on "the general intent of Congress, the purposes of the patent system, and the implications of a decision one way or the other." Id. at 532, 148 USPQ at 695. The Court concluded that "[t]he basic quid pro quo contemplated by the Constitution and the Congress for granting a patent monopoly is the benefit derived by the public from an invention with substantial utility. Unless and until *1024 a process is refined and developed to this point--where specific benefit exists in currently available form--there is insufficient justification for permitting an applicant to engross what may prove to be a broad field." Id. at 534-35, 148 USPQ at 695.

The Court considered and rejected the applicant's argument that attenuating the requirement of utility "would encourage inventors of new processes to publicize the event for the benefit of the entire scientific community, thus widening the search for uses and increasing the fund of scientific knowledge." The Court noted that, while there is value to encouraging disclosure, "a more compelling consideration is that a process patent in the chemical field, which has not been developed and pointed to the degree of specific utility, creates a monopoly of knowledge which should be granted only if clearly commanded by the statute. Until the process claim has been reduced to production of a product shown to be useful, the metes and bounds of that monopoly are not capable of precise delineation. It may engross a vast, unknown, and perhaps unknowable area. Such a patent may confer power to block off whole areas of scientific development." Id. at 534, 148 USPQ at 695.

The Court took pains to note that it did not "mean to disparage the importance of contributions to the fund of scientific information short of the invention of something 'useful,'" and that it was not "blind to the prospect that what now seems without 'use' may tomorrow command the grateful attention of the public." Id. at 535-36, 148 USPQ at 696. Those considerations did not sway the Court, however, because "a patent is not a hunting license. It is not a reward for the search, but compensation for its successful conclusion." Id.

Subsequent decisions of the CCPA and the Court of Appeals for the Federal Circuit have added further layers of judicial gloss to the meaning of § 101's utility requirement. The first opinion of the CCPA applying *Brenner* was *In re Kirk*, 376 F.2d 936, 153 USPQ 48 (CCPA 1967). The invention claimed in *Kirk* was a set of steroid derivatives said to have valuable biological properties and to be of value "in the furtherance of steroidal research and in the application of steroidal materials to veterinary or medical practice." Id. at 938, 153 USPQ at 50. The claims had been rejected for lack of utility. In response, the applicants submitted an affidavit which purportedly "show[ed] that one skilled in the art would be able to determine the biological uses of the claimed compounds by routine tests." Id. at 939, 153 USPQ at 51.

The court held that "nebulous expressions [like] 'biological activity' or 'biological properties'" did not adequately convey how to use the claimed compounds. Id. at 941, 153 USPQ at 52. Nor did the applicants' affidavit help their case: "the sum and substance of the affidavit appear[ed] to be that one of ordinary skill in the art would know 'how to use' the compounds to find out in the first instance whether the compounds are--or are not--in fact useful or possess useful properties, and to ascertain what those properties are." Id. at 942, 153 USPQ at 53.

The *Kirk* court held that an earlier CCPA decision, holding that a chemical compound meets the requirements of § 101 if it is useful to chemists doing research on steroids, had effectively been overruled by *Brenner*. "There can be no doubt that the insubstantial, superficial nature of vague, general disclosures or arguments of 'useful in research' or 'useful as building blocks of value to the researcher' was recognized, and clearly rejected, by the Supreme Court" in *Brenner*. See *Kirk*, 376 F.2d at 945, 153 USPQ at 55.

More recently, in *In re Ziegler*, 992 F.2d 1197, 26 USPQ2d 1600 (Fed. Cir. 1993), the Federal Circuit considered the degree of specificity required to show utility for a claim to polypropylene. The U.S. application on appeal in *Ziegler* claimed priority to a German application filed in 1954. "In the German application, Ziegler disclosed only that solid granules of polypropylene could be pressed into a flexible film with a characteristic infrared spectrum and that the polypropylene was 'plastic-like.'" Id. at 1203, 26 USPQ2d at 1605. "Ziegler did not assert any practical use for the polypropylene or its film, and Ziegler did not disclose any characteristics of the polypropylene

or its film that demonstrated its utility." *Id.* The court held that the German application did not satisfy the requirements of § 101 and therefore could not be relied on to overcome a rejection based on an intervening reference. *Id.* "[At] best, Ziegler was on the way to discovering a practical utility for polypropylene at the time of the filing of the German application; but in that application Ziegler had not yet gotten there." *Id.*

On the other hand, the CCPA reversed a rejection for lack of utility in *In re Jolles*, 628 F.2d 1322, 206 USPQ 885 (CCPA 1980). The applicant in *Jolles* claimed pharmaceutical compositions that were disclosed to be useful in treating acute myeloblastic leukemia. *1025 *See id.* at 1323, 206 USPQ at 886. The active ingredients in the compositions were closely related to daunorubicin and doxorubicin, both of which were "well recognized in the art as valuable for use in cancer chemotherapy." *Id.*, 206 USPQ at 887. The applicant also submitted declaratory evidence showing that eight of the claimed compositions were effective in treating tumors in a mouse model, and one was effective in treating humans. *See id.* at 1323-24, 206 USPQ at 887-88. The court noted that the data derived from the mouse model were "relevant to the treatment of humans and [were] not to be disregarded," *id.* at 1327, 206 USPQ at 890, and held that the evidence was sufficient to support the asserted therapeutic utility. *See id.* at 1327-28, 206 USPQ at 891.

The Federal Circuit held in *Cross v. Iizuka*, 753 F.2d 1040, 224 USPQ 739 (Fed. Cir. 1985), that *in vivo* testing (as in *Jolles*) was not necessarily required to show utility in the pharmaceutical context. The *Cross* court stated that "[it] is axiomatic that an invention cannot be considered 'useful,' in the sense that a patent can be granted on it, unless substantial or practical utility for the invention has been discovered and disclosed where such utility would not be obvious." *Id.* at 1044, 224 USPQ at 742 (citing *Brenner v. Manson*). The court "perceive[d] no insurmountable difficulty, under appropriate circumstances, in finding that the first link in the screening chain, *in vitro* testing, may establish a practical utility for the compound in question." *Id.* at 1051, 224 USPQ at 748. Successful *in vitro* testing could provide an immediate benefit to the public, by "marshall[ing] resources and direct[ing] the expenditure of effort to further *in vivo* testing of the most potent compounds . . . , analogous to the benefit provided by the showing of an *in vivo* utility." *Id.* On the facts of that case -- successful *in vitro* testing supplemented by similar *in vitro* and *in vivo* activities of structurally similar compounds -- the court held

that *in vitro* activity was sufficient to meet the requirements of § 101. *See id.*

The Federal Circuit confirmed in *In re Brana*, 51 F.3d 1560, 34 USPQ2d 1436 (Fed. Cir. 1995), that human testing is not necessary to establish utility for a method of treatment. The invention claimed in *Brana* was a group of compounds disclosed to have antitumor activity. *See id.* at 1562, 34 USPQ2d at 1437-38. The specification disclosed that the claimed compounds had higher antitumor activity than related compounds known to have antitumor activity, and the applicants provided declaratory evidence of *in vivo* activity against tumors in a mouse model. *See id.*, 34 USPQ2d at 1438. The court held that these data were sufficient to satisfy § 101; usefulness in patent law does not require that the invention be ready to be administered to humans. *See id.* at 1567, 34 USPQ2d at 1442.

[1] Several lessons can be drawn from *Brenner* and its progeny. First, § 101's requirement that an invention be "useful" is not to be given its broadest reach, such that little or nothing of a chemical nature would be found to lack utility. *See Brenner*, 383 U.S. at 530, 148 USPQ at 694. Thus, not every "use" that can be asserted will be sufficient to satisfy § 101. For example, the steroid compound at issue in *Brenner* was useful as a possible object of scientific inquiry, and the polypropylene claimed in *Ziegler* was useful for pressing into a flexible film, yet both lacked sufficient utility to satisfy § 101. *See Brenner*, 383 U.S. at 529, 148 USPQ at 696; *Ziegler*, 992 F.2d at 1203, 26 USPQ2d at 1605.

Rather than setting a *de minimis* standard, § 101 requires a utility that is "substantial", i.e., one that provides a specific benefit in currently available form. *Brenner*, 383 U.S. at 534-35, 148 USPQ at 695. This standard has been found to be met by pharmaceutical compositions shown to be useful in mouse models and in humans for treating acute myeloblastic leukemia (*Jolles*, 628 F.2d at 1327-28, 206 USPQ at 891); by evidence showing successful *in vitro* testing supplemented by similar *in vitro* and *in vivo* activities of structurally similar compounds (*Cross*, 753 F.2d at 1051, 224 USPQ at 748); and by evidence showing *in vivo* antitumor activity in mice, combined with a disclosure that the claimed compounds had higher antitumor activity than a related compound known to have antitumor activity (*Brana*, 51 F.3d at 1567, 34 USPQ2d at 1442).

By contrast, *Brenner's* standard has been interpreted to mean that "vague, general disclosures or

arguments of 'useful in research' or 'useful as building blocks of value to the researcher'" would not satisfy § 101. See *Kirk*, 376 F.2d at 945, 153 USPQ at 55 (interpreting *Brenner*). Likewise, a disclosure of a "plastic-like" polypropylene capable of being pressed into a flexible film was held to show that the applicant was "at best . . . on the way to discovering a practical utility for polypropylene at the time of the filing," but not yet there. *Ziegler*, at 1203, 26 USPQ2d at 1605.

With these principles in mind we turn to the issues at hand. Of the many utilities asserted *1026 in the specification, two have received the most attention in the briefing in this appeal, *i.e.*, identification and detection of polymorphisms and use as probes or as a source for primers. We shall focus on these asserted utilities first and then address the other arguments set forth in the briefing.

a. Polymorphisms

[2] This utility is discussed at pages 35-42 of the specification in terms of what polymorphisms are and how one would go about determining the existence of a polymorphism. The discussion in this portion of the specification, however, is not specific to the nucleotide molecules depicted in SEQ ID NO: 1 through SEQ ID NO: 5. To the contrary, according to appellants' specification (page 35, lines 25-26), "one or more of the [32,236] EST nucleic acid molecules (or a sub-fragment thereof) may be employed as a marker nucleic acid molecule to identify . . . polymorphism(s)." The specification does not explain why any of the 32,236 nucleotide molecules disclosed in the specification, or more specifically the five nucleotide molecules depicted in SEQ ID NO: 1 through SEQ ID NO: 5, would in fact be useful in detecting polymorphisms.

Rather, appellants argue (Brief, page 7), "the claimed nucleic acid molecules have utility even if the absence of a particular polymorphism is detected. Indeed, the absence of a polymorphism usually demonstrates that the two (or more) populations being compared share a common genetic heritage." In other words, appellants' position is that an EST by definition possesses patentable utility because it can be used by itself in determining whether populations share a common genetic heritage. While that may be a "utility," we do not find that it is a *substantial* utility.

Without knowing any further information in regard to the gene represented by an EST, as here, detection of the presence or absence of a polymorphism

provides the barest information in regard to genetic heritage. As the examiner explains (Answer, bridging paragraph, pages 10-11):

Polymorphisms are natural variations within sequences which themselves may not have any meaningful use. Therefore, determining whether the claimed nucleic acids [(or nucleic acids detected by the claimed nucleic acids)] have or do not have a polymorphism would require determining whether there was a polymorphism within such a sequence and then determining how to use this information in a patentably meaningful way. The [a]ppellant also argues, "many of these uses are directly analogous to a microscope". This argument has been reviewed but is not convincing because the microscope provides information to the scientist which is automatically useful. For example, the microscope may be used for identification and differentiation between gram-positive and gram-negative bacteria. The differentiation of bacteria facilitates in the administration of proper antibiotics. For example, if the microscope is used to determine whether Staph is present or whether Strep is present provides valuable information to the scientist and/or doctor for treating patients. The instant invention, however, provides no information to this extent. If the scientist determines that SEQ ID NO: 1 is present, the scientist does not know how to use this information. Thus, the identification of SEQ ID NO: 1 is not a substantial utility.

In contrast, at the other end of the "utility spectrum" would be information gleaned from detecting the presence or absence of a polymorphism when it is known what effect the gene from which the EST is derived has in the development and/or phenotype of the plant. Somewhere between having no knowledge (the present circumstances) and having complete knowledge of the gene and its role in the plant's development and/or phenotype lies the line between "utility" and "substantial utility." We need not draw the line or further define it in this case because the facts in this case represent the lowest end of the spectrum, *i.e.*, an insubstantial use.

b. Probes or source of primers

[3] Appellants argue that the "specification discloses that the claimed nucleic acid molecules can be used to isolate nucleic acid molecules of other plants and organisms. . . ." Appeal Brief, page 8. While that may be true, it begs the question of what substantial use such nucleic acid molecules would have? Again, the present specification does not attribute any property in terms of plant trait, or phenotype to any of the nucleotide molecules set

forth in SEQ ID NO: 1 through SEQ ID NO: 5. In the absence of such information, using the claimed molecules to isolate other molecules, which themselves lack substantial utility, does not represent a substantial utility.

Appellants also assert that the claimed nucleic acid molecules may be used in a *1027 "chromosome walk." Brief, pages 8-9. According to appellants (Brief, page 9),

The claimed nucleic acid molecules provide a particularly appropriate and demonstrably useful starting point for a walk to isolate a promoter that is active in leaves at the time of anthesis. Isolation of such a promoter would be desirable and particularly useful because it allows expression of proteins at that important developmental state, including proteins that provide disease resistance. Because the claimed nucleic acid molecules were isolated from leaves, they provide an appropriate starting point for isolating a promoter active in leaves. A random nucleic acid molecule does not provide an equally good starting point to isolate such a promoter.

As we understand this argument, the claimed ESTs may be useful in searching for promoters that are only active in leaves at the time of anthesis. The specification, however, fails to demonstrate that any of the nucleic acid molecules set forth in SEQ ID NO: 1 through SEQ ID NO: 5 would be useful in obtaining a successful result from such a search. As set forth at page 34, lines 14-19 of appellants' specification,

The [32,236] nucleic acid molecules of the present invention may be used to isolate promoters of tissue enhanced[,] tissue specific, cell-specific, cell-type, developmentally or environmentally regulated expression profiles. Isolation and functional analysis of the 5' flanking promoter sequences of these genes from genomic libraries, for example, using genomic screening methods and PCR techniques would result in the isolation of useful promoters and transcriptional regulatory elements.

The specification does not provide any expectation of successfully using any of the 32,236 nucleic acid molecules disclosed in the specification, or more specifically the five nucleic acid molecules depicted in SEQ ID NO: 1 through SEQ ID NO: 5, to isolate promoters of tissue enhanced, tissue specific, cell-specific, cell-type, developmentally or environmentally regulated expression profiles.

Furthermore, notwithstanding appellants' assertion

(Brief, page 9), there is no evidence on this record that any of the nucleic acid molecules depicted in SEQ ID NO: 1 through SEQ ID NO: 5 are tissue or cell-type specific, or developmentally or environmentally regulated. In this regard, we note that the claimed nucleic acid molecules were isolated from the cDNA library LIB3115. Specification, page 80, lines 5-6. There is no evidence on this record that LIB3115 is a subtractive cDNA library, wherein nucleic acid molecules from other maize tissue, or from other developmental stages, was subtracted (removed) from the library. Compare, for example, the subtractive cDNA library LIB3153 which is disclosed (specification, page 83, lines 17-19) to be "generated by subtracting driver cDNA, which is prepared from kernels harvested from 15 DAP [days after pollination] maize plants, from target cDNA, which is prepared from endosperms harvested from 5-8 day[s] after pollination (DAP) maize plants." In contrast to the claimed nucleic acid molecules, nucleic acid molecules SEQ ID NO: 24,931 through SEQ ID NO: 25,680 are from the subtractive cDNA library LIB3153.

In our opinion, the claimed nucleic acid molecules having the sequences identified as SEQ ID NO: 1 through SEQ ID NO: 5, represent five randomly selected nucleic acid molecules isolated from pooled leaf tissue at the time of anthesis. Notwithstanding appellants' emphasis on "anthesis," for the foregoing reasons, we find no evidence on this record that any of appellants' five randomly selected nucleic acid molecules are expressed only at the time of "anthesis." Accordingly, despite appellants' assertion to the contrary, there is no reasonable expectation that any of the claimed nucleic acid molecules would be capable of isolating a promoter that was only active in leaves at the time of anthesis. As appellants recognize (Brief, page 9), "[a] random nucleic acid molecule does not provide an equally good starting point to isolate such a promoter" compared to a nucleic acid molecule that is known to be specifically associated with this stage of plant development.

We recognize appellants' argument (Brief, bridging sentence, pages 9-10), "[a]n invention may be 'less effective than existing devices but nevertheless meet the statutory criteria for patentability.' Custom Accessories, Inc. v. Jeffrey-Allan Indus., 807 F.2d 955, 960 n.12, 1 U.S.P.Q.2d 1196, 1199 n.12 (Fed. Cir. 1986)." While we agree with appellants' statement, we fail to see how it applies to appellants' claimed invention, wherein there is no evidence or expectation that the claimed nucleic acid molecules would be "effective" at all. In this regard, we remind appellants that an invention does not have utility

sufficient to satisfy § 101 until it is "refined and developed" to the point of providing a specific benefit *1028 in currently available form. *See, e.g., Brenner*, 383 U.S. at 534, 148 USPQ at 695.

[4] An invention certainly can have a utility that is shared by other compounds or compositions. Take, for example, an application that claims ibuprofen and discloses that it is useful as an analgesic. No one would argue that a claim to ibuprofen lacks utility simply because aspirin and acetaminophen are also useful as analgesics. On the other hand, not every utility will satisfy § 101, even if the utility is shared by a class of inventions. Assume that the above-described application did not disclose that ibuprofen was an analgesic but only disclosed that it is useful because it can be used to fill a jar, which would then be useful as a paperweight. There would be little doubt that this disclosed utility would not satisfy § 101, even though the utility is shared by a large class of inventions, viz., those whose physical embodiments have mass. So while a utility need not be unique to a claimed invention, it must nonetheless be specific, and in currently available form, in order to satisfy § 101.

c. Other Arguments

Appellants argue that the specification "discloses additional utilities for the claimed nucleic acid molecules including introduction of the claimed nucleic acid molecules into a plant or plant cell (either as sense or antisense inhibitors), which can then be used to screen for compounds such as a herbicide." Brief, page 6. Specifically, appellants argue (*id.*) that "a compound can be provided to both an antisense plant and a control plant (no antisense) and the effect of the compound on the plant can be monitored." Appellants analogize this proposed procedure to a "cell-based assay" which appellants assert to have a "legally sufficient utility." *Id.*

[5] Suffice it to say that an otherwise uncharacterized nucleic acid molecule is being claimed in this application, not an assay. The portion of the specification cited in support of this argument (page 73, line 17 through page 74, line 17) indicates that the nucleic acid molecule must be introduced into a plant cell and transcribed using an appropriate promoter to result in the suppression of an endogenous protein. The specification does not indicate that such a method is feasible when the nucleic acid to be used is uncharacterized [FN5] as here. Such a use does not provide a specific or substantial benefit in currently available form.

[6] Appellants also argue that the claimed nucleic acids are useful to measure the level of mRNA in a sample through use of microarray technology and use as molecular markers. Brief, page 6. In regard to microarrays, appellants argue (*id. fn. 3*) that it is "standard practice" to screen populations of nucleic acids with EST sequences without characterizing each and every target mRNA. We find that the asserted utility of the claimed nucleic acid--as one component of an assay for monitoring gene expression--does not satisfy the utility requirement of § 101. Such a use does not provide a specific benefit in currently available form. We accept, for argument's sake, that a person skilled in the art could use the claimed nucleic acid, in combination with other nucleic acids, to monitor changes in expression of the gene that encompasses the nucleic acid depicted in e.g., SEQ ID NO: 1. However, the specification provides no guidance that would allow a skilled artisan to use data relating to expression of such a gene in any practical way. The specification simply provides no guidance regarding what the SEQ ID NO: 1-specific information derived from a gene expression experiment would mean. As the examiner points out (Answer, page 9), "the instant claimed nucleic acids appear to require further experimentation on the material itself to determine the function and properties of the claimed nucleic acids."

To highlight the examiner's assertion, suppose, for example, that a researcher found that SEQ ID NO: 1 expression was increased when a cell was treated with a particular agent. The specification provides no basis on which a skilled worker would be able to determine whether that result is meaningful. Maybe the meaning in a change in SEQ ID NO: 1 expression would depend on other factors, but again the specification provides no hint as to what other factors might be important. Would it depend on what agent is used, what cell type is used, the behavior of other genes (if so, which genes and what behavior is significant), the degree of increase? The specification simply provides no guidance as to how to interpret the results that might be seen using SEQ ID NO: 1 in a gene expression assay.

*1029 In effect, appellants' position is that the claimed nucleic acids are useful because those of skill in the art could experiment with them and figure out for themselves what any observed experimental results might mean. We do not agree that such a disclosure provides a "specific benefit in currently available form." Rather, the present case seems analogous to *Brenner*. In *Brenner*, the applicant

claimed a method of making a compound but disclosed no utility for the compound. 383 U.S. at 529, 148 USPQ at 693. The Court held that a process lacks utility if it produces a product that lacks utility. Id. at 534, 148 USPQ at 695. Here, appellants claim a product asserted to be useful in a method of generating gene-expression data, but the specification does not disclose how to interpret those data. Just as the process claimed in *Brenner* lacked utility because the specification did not disclose how to use the end-product, the products claimed here lack utility, because even if used in gene expression assays, the specification does not disclose how to use SEQ ID NO: 1-specific gene expression data.

Assuming *arguendo*, that a generic gene expression assay--one based on monitoring expression of thousands of uncharacterized nucleic acids would provide a useful tool for, e.g., drug discovery, it does not follow that each one of the nucleic acids represented in the assay individually has patentable utility. Although each nucleic acid in the assay contributes to the data generated by the assay overall, the contribution of a single nucleic acid--its data point--is only a tiny contribution to the overall picture. The *Brenner* Court held that § 101 sets more than a *de minimis* standard for utility. Therefore, the patentable utility of a gene expression assay, for example, does not necessarily mean that each tiny component of the assay also has patentable utility. A patentable utility divided by a thousand does not necessarily equal a thousand patentable utilities. Each claimed invention must be shown to meet § 101's utility requirement in order to be patentable; it must provide a specific benefit in currently available form. Providing a single data point among thousands or millions, even if the thousands or millions of data points collectively are useful, does not meet this standard. The Supreme Court noted that the patent system contemplates a basic quid pro quo: in exchange for the legal right to exclude others from his invention for a period of time, an inventor discloses his invention to the public. See *Brenner*, 383 U.S. at 534, 148 USPQ at 695. The *Brenner* Court held that the grant of patent rights to an applicant is justified only by disclosure of an invention with substantial utility -- a specific benefit in currently available form. Until the invention has been refined and developed to this point, the Court held, the applicant has not met his side of the bargain, and has not provided a disclosure sufficient to justify a grant of the right to exclude others. See *id.*

We reach the same conclusion in regard to appellants' assertion that the nucleic acid molecules depicted in SEQ ID NO: 1 through SEQ ID NO: 5 are

useful as a molecular marker or probe. It is not seen that the one data point which may be provided by using the uncharacterized nucleic acid molecule of SEQ ID NO: 1 as a molecular marker or probe represents a substantial use.

Appellants argue that ESTs have real world value as seen from the "growth of a multi-million dollar industry in the United States premised on the usefulness of ESTs." Brief, page 11. Since appellants fail to provide any suggestion on which use of ESTs this industry is premised on, we can only assume that appellants are referring to the potential usefulness of EST databases, clone sets or microarrays. Suffice it to say, the claims on appeal are not directed to EST databases, clone sets and/or microarrays. Again, it is not seen that the one data point which may be provided by using the uncharacterized nucleic acid molecules of SEQ ID NO: 1 through SEQ ID NO: 5 in such devices represents a substantial use.

For the foregoing reasons we affirm the rejection of claim 1 under 35 U.S.C. § 101.

Enablement

According to the examiner (Answer, page 13, emphasis removed), "since the claimed invention is not supported by either a specific, substantial asserted utility or a well established utility for the reasons set forth [in support of the rejection under 35 U.S.C. § 101] one skilled in the art clearly would not know how to use the claimed invention." This rejection is simply a corollary of the finding of lack of utility. Appellants assert (Brief, page 12), this rejection should be reversed for the same reasons set forth in their arguments regarding the rejection under 35 U.S.C. § 101. Thus, our conclusion with respect to the § 101 issue will also apply to this aspect of the § 112 (enablement) issue. On this basis we affirm the rejection of claim 1 under the enablement provision of 35 U.S.C. § 112, first paragraph.

*1030 Written description

This rejection stands on a different footing. As we understand the examiner's argument the use of the transitional phrase "comprising" in appellants' claimed invention results in appellants claiming a large genus of nucleic acid molecules which are not adequately described by SEQ ID NO: 1 through SEQ ID NO: 5. Answer, pages 13-16. Apparently the examiner is of the opinion that the claimed invention should be limited to nucleic acid molecules as set forth in SEQ ID NO: 1 through SEQ ID NO: 5. In response appellants argue (Brief, page 14, original footnote omitted),

Applicants have provided the nucleotide sequences required by the claims, *i.e.*, SEQ ID NO: 1, SEQ ID NO: 2, SEQ ID NO: 3, SEQ ID NO: 4, and SEQ ID NO: 5, and have thus established possession of the claimed invention. The fact that the claims at issue are intended to cover molecules that include the recited sequences joined with additional sequences [FN6]] does not mean that [a]pplicants were any less in possession of the claimed nucleic acid molecules.

[7] As discussed *supra*, as we understand the claimed invention, the use of the transitional term "comprising" does not allow for internal alterations (e.g. insertions or deletions) of the nucleotide sequences set forth in SEQ ID NO: 1 through SEQ ID NO: 5, but instead only allows for the addition of nucleotides or other molecules at either end of the nucleotide sequences set forth in SEQ ID NO: 1 through SEQ ID NO: 5. We agree with appellants that they have provided an adequate written description of nucleic acid molecules with the sequences set forth in SEQ ID NO: 1 through SEQ ID NO: 5. That the claimed nucleic acid molecules may have other molecules attached to either, or both of their 5' or 3' ends does not diminish appellants' adequate written description of the nucleic acids molecules with the sequences set forth in SEQ ID NO: 1 through SEQ ID NO: 5 as claimed.

Accordingly, we reverse the rejection of claim 1 under the written description provision of 35 U.S.C. § 112, first paragraph.

No time period for taking any subsequent action in connection with this appeal may be extended under 37 CFR § 1.136(a).

AFFIRMED

FN1. During the Oral Hearing, appellants' representative confirmed that the administrative file contained no evidence that the claimed ESTs were capable of detecting a polymorphism that correlated with any particular trait.

FN2. This interpretation of the claimed invention was confirmed by appellants' representative during the Oral Hearing.

FN3. In discussing the issue of utility under 35 U.S.C. § 101, the Federal Circuit and the Court of Customs and Patent Appeals since *Brenner*, have used the phrases "substantial utility" and "practical utility"

interchangeably. See *e.g.*, *Fujikawa v. Wattanasin*, 93 F.3d 1559, 1963-1964, 39 USPQ2d 1895, 1898-1899 (Fed. Cir. 1996) ("It is well established that a patent may not be granted to an invention unless substantial or practical utility for the invention has been discovered and disclosed.").

FN4. The invention at issue in *Brenner* was a process, but the Court expressly noted that its holding "would apply equally to the patenting of the product produced by the process." *Id.* at 535, 148 USPQ at 695-96.

FN5. To emphasize the uncharacterized nature of appellants' invention we note the examiner's finding (Answer, page 17) that translating SEQ ID NO: 5 in all 6 possible reading frames reveals that the sequence contains numerous stop codons which would terminate the translation of a protein, or protein fragment, encoded thereby.

FN6. By way of examples appellants explain (Brief, bridging paragraph, pages 14-15) that the specification discloses, *inter alia*, the claimed nucleic acid molecules joined together with vectors, and other nucleic acids (e.g. fusion nucleic acid molecules) and detectable labels.

982 F.2d 1554, 1560 (Fed.Cir.1992) (holding a C.F.R. provision invalid because it did not "comport with the clear statutory mandate"). Because § 831.2203(f) conflicts with and is no longer supported by the amendments to § 8343a, it is ineffective. Resort to that regulation to affect an AFA following the effective dates of amendments is foreclosed.

Here, Kievenaar's husband completed the annuity paperwork in February 2002. His annuity was to begin on July 1, 2002, and he died suddenly on July 22, 2002. Thus, his annuity commenced at a time when § 831.2203(f) was ineffective. While we are sympathetic to Kievenaar's position, we have no alternative but to conclude that she is not eligible for the AFA she seeks.

Because the unavailability of § 831.2203(f) during the period relevant to Kievenaar is dispositive of her case, we need not and do not decide the issue of validity of the regulation prior to the effective dates of the amendments to § 8343a.

CONCLUSION

The Board correctly determined that 5 C.F.R. § 831.2203(f) was inapplicable to Kievenaar's request to change her husband's self-only annuity to an AFA. The Board's affirmance of OPM's decision to deny the requested AFA was in accordance with law and is affirmed.

AFFIRMED



**In re Dane K. FISHER and
Raghunath V. Lalgudi.**

No. 04-1465.

United States Court of Appeals,
Federal Circuit.

Sept. 7, 2005.

Background: Applicants for patent covering "expressed sequence tags" for identifying nucleic acid sequences in maize genes challenged decision of United States Patent and Trademark Office Board of Patent Appeals and Interferences, 72 U.S.P.Q.2d 1020, 2004 WL 2185929, finding their claim unpatentable for lack of utility and lack of enablement.

Holdings: The Court of Appeals, Michel, Chief Judge, held that:

- (1) claimed invention lacked specific and substantial utility, and
- (2) application failed for lack of enablement.

Affirmed.

Rader, Circuit Judge, dissented and filed opinion.

1. Patents ⇨314(5)

Whether patent application discloses requisite utility for claimed invention is question of fact. 35 U.S.C.A. § 101.

2. Patents ⇨48

Patent application covering "expressed sequence tags" for identifying nucleic acid sequences in maize genes lacked requisite utility where tagged genes had no known functions; claimed tags acted as no more than research intermediates that might help scientists to isolate particular underlying protein-encoding genes and conduct further experimentation on those genes. 35 U.S.C.A. § 101.

3. Patents ⇌46

To be patentable, claimed invention must have substantial and specific "utility," i.e., invention must have significant and presently available benefit to public, and its claimed use must not be so vague as to be meaningless. 35 U.S.C.A. § 101.

See publication Words and Phrases for other judicial constructions and definitions.

4. Evidence ⇌48

Patents ⇌97

Manual of Patent Examining Procedure (MPEP) and Patent and Trademark Office's Utility Guidelines are not binding on court ruling on patent's validity, but may be given judicial notice to extent they do not conflict with patent statutes.

5. Patents ⇌99

Patent application for invention that lacks requisite utility also fails, as matter of law, for lack of enablement. 35 U.S.C.A. §§ 101, 112(1).

Seth P. Waxman, Wilmer Cutler Pickering Hale and Dorr LLP, of Washington, DC, argued for appellants. With him on the brief were William F. Lee and Richard W. O'Neill, of Boston, Massachusetts; and William G. McElwain and Henry N. Wixon, of Washington, DC.

Stephen Walsh, Associate Solicitor, United States Patent and Trademark Office, of Arlington, Virginia, argued for the Director of the Patent and Trademark Office. With him on the brief were John M. Whealan, Solicitor, and Thomas W. Krause, Associate Solicitor.

Joseph A. Keyes, Jr., of Washington, DC, for amicus curiae Association of American Medical Colleges.

1. The real party in interest is Monsanto Technology LLC, which is owned by the Monsanto

Marc S. Gold, of Washington, DC, for amicus curiae National Academy of Sciences.

Donald R. Stuart, of Indianapolis, Indiana, for amicus curiae Dow AgroSciences LLC. With him on the brief was Kenneth B. Ludwig.

Paula K. Davis, of Indianapolis, Indiana, for amicus curiae Eli Lilly and Company. With her on the brief were Steven P. Caltrider and James J. Kelley.

Michael C. Schiffer, of Irvine, California, for amicus curiae Baxter Healthcare Corporation.

Darrel C. Karl, Finnegan, Henderson, Farabow, Garrett & Dunner, L.L.P., of Washington, DC, for amicus curiae American College of Medical Genetics.

Jeffrey P. Kushan, Sidley Austin Brown & Wood, LLP, of Washington, DC, for amicus curiae Genentech, Inc. With him on the brief were Kathi A. Cover and David L. Fitzgerald.

George C. Yu, of Emeryville, California, for amicus curiae Affymetrix, Inc.

Before MICHEL, Chief Judge, RADER and BRYSON, Circuit Judges.

Opinion for the court filed by Chief Judge MICHEL.

Dissenting opinion filed by Circuit Judge RADER.

MICHEL, Chief Judge.

Dane K. Fisher and Raghunath Lalgudi (collectively "Fisher")¹ appeal from the decision of the U.S. Patent and Trademark Office ("PTO") Board of Patent Appeals and Interferences ("Board") affirming the examiner's final rejection of the only pending claim of application Serial No. 09/619,-

Company.

643 (the "'643 application"), entitled "Nucleic Acid Molecules and Other Molecules Associated with Plants," as unpatentable for lack of utility under 35 U.S.C. § 101 and lack of enablement under 35 U.S.C. § 112, first paragraph. *Ex parte Fisher*, App. No.2002-2046 (Bd.Pat.App.Int. Mar. 16, 2004) ("Board Decision"). This appeal was submitted after oral argument on May 3, 2005. Because we conclude that substantial evidence supports the Board's findings that the claimed invention lacks a specific and substantial utility and that the '643 application does not enable one of ordinary skill in the art to use the invention, we affirm.

I. BACKGROUND

A. Molecular Genetics and ESTs

The claimed invention relates to five purified nucleic acid sequences that encode proteins and protein fragments in maize plants. The claimed sequences are commonly referred to as "expressed sequence tags" or "ESTs." Before delving into the specifics of this case, it is important to understand more about the basic principles of molecular genetics and the role of ESTs.

Genes are located on chromosomes in the nucleus of a cell and are made of deoxyribonucleic acid ("DNA"). DNA is composed of two strands of nucleotides in double helix formation. The nucleotides contain one of four bases, adenine ("A"), guanine ("G"), cytosine ("C"), and thymine ("T"), that are linked by hydrogen bonds to form complementary base pairs (*i.e.*, A-T and G-C).

When a gene is expressed in a cell, the relevant double-stranded DNA sequence is transcribed into a single strand of messenger ribonucleic acid ("mRNA"). Messen-

ger RNA contains three of the same bases as DNA (A, G, and C), but contains uracil ("U") instead of thymine. mRNA is released from the nucleus of a cell and used by ribosomes found in the cytoplasm to produce proteins.

Complementary DNA ("cDNA") is produced synthetically by reverse transcribing mRNA. cDNA, like naturally occurring DNA, is composed of nucleotides containing the four nitrogenous bases, A, T, G, and C. Scientists routinely compile cDNA into libraries to study the kinds of genes expressed in a certain tissue at a particular point in time. One of the goals of this research is to learn what genes and downstream proteins are expressed in a cell so as to regulate gene expression and control protein synthesis.²

An EST is a short nucleotide sequence that represents a fragment of a cDNA clone. It is typically generated by isolating a cDNA clone and sequencing a small number of nucleotides located at the end of one of the two cDNA strands. When an EST is introduced into a sample containing a mixture of DNA, the EST may hybridize with a portion of DNA. Such binding shows that the gene corresponding to the EST was being expressed at the time of mRNA extraction.

Claim 1 of the '643 application recites:

A substantially purified nucleic acid molecule that encodes a maize protein or fragment thereof comprising a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 5.

The ESTs set forth in SEQ ID NO: 1 through SEQ ID NO: 5 are obtained from cDNA library LIB3115, which was generated from pooled leaf tissue harvested

2. We have discussed the basic principles of molecular genetics more extensively in prior cases. See, *e.g.*, *In re Deuel*, 51 F.3d 1552, 1554-56 (Fed.Cir.1995); *Amgen, Inc. v. Chu-*

gai Pharm. Co., Ltd., 927 F.2d 1200, 1207-08 (Fed.Cir.1991); *In re O'Farrell*, 853 F.2d 894, 895-99 (Fed.Cir.1988).

from maize plants (RX601, Asgrow Seed Company, Des Moines, Iowa, U.S.A.) grown in the fields at Asgrow research stations. SEQ ID NO:1 through SEQ ID NO:5 consist of 429, 423, 365, 411, and 331 nucleotides, respectively. When Fisher filed the '643 application, he claimed ESTs corresponding to genes expressed from the maize pooled leaf tissue at the time of anthesis. Nevertheless, Fisher did not know the precise structure or function of either the genes or the proteins encoded for by those genes.

The '643 application generally discloses that the five claimed ESTs may be used in a variety of ways, including: (1) serving as a molecular marker for mapping the entire maize genome, which consists of ten chromosomes that collectively encompass roughly 50,000 genes; (2) measuring the level of mRNA in a tissue sample via microarray technology to provide information about gene expression; (3) providing a source for primers for use in the polymerase chain reaction ("PCR") process to enable rapid and inexpensive duplication of specific genes; (4) identifying the presence or absence of a polymorphism; (5) isolating promoters via chromosome walking; (6) controlling protein expression; and (7) locating genetic molecules of other plants and organisms.

B. Final Rejection

In a final rejection, dated September 6, 2001, the examiner rejected claim 1 for lack of utility under § 101. The examiner found that the claimed ESTs were not supported by a specific and substantial utility. She concluded that the disclosed uses were not specific to the claimed ESTs, but instead were generally applicable to any EST. For example, the examiner noted that any EST may serve as a molecular tag to isolate genetic regions. She also concluded that the claimed ESTs lacked a substantial utility because there was no known use for the proteins pro-

duced as final products resulting from processes involving the claimed ESTs. The examiner stated: "Utilities that require or constitute carrying out further research to identify or reasonably confirm a 'real world' context of use are not substantial utilities."

The examiner also rejected the claimed application for lack of enablement under § 112, first paragraph. She reasoned that one skilled in the art would not know how to use the claimed ESTs because the '643 application did not disclose a specific and substantial utility for them.

On July 19, 2000, Fisher filed a notice of appeal with the Board.

C. Board Proceedings

The Board considered each of Fisher's seven potential uses but noted that Fisher focused its appeal on only two: (1) use for the identification of polymorphisms; and (2) use as probes or as a source for primers. As to the first, the Board found that the application failed to explain why the claimed ESTs would be useful in detecting polymorphisms in maize plants. *Board Decision*, slip op. at 14. The Board reasoned that "[w]ithout knowing any further information in regard to the gene represented by an EST, as here, detection of the presence or absence of a polymorphism provides the barest information in regard to genetic heritage." *Id.*, slip op. at 15. Thus, the Board concluded that Fisher's asserted uses for the claimed ESTs tended to the "insubstantial use" end of the spectrum between a substantial and an insubstantial utility. *Id.*

The Board also concluded that using the claimed ESTs to isolate nucleic acid molecules of other plants and organisms, which themselves had no known utility, is not a substantial utility. *Id.*, slip op. at 16. Specifically, the Board noted that Fisher argued that the "claimed ESTs may be

useful in searching for promoters that are only active in leaves at the time of anthesis.” *Id.* The Board found, however, that the application failed to show that the claimed ESTs would be expressed only during anthesis or that they would be capable of isolating a promoter active in maize leaves at the time of anthesis. *Id.*, slip op. at 18.

Additionally, the Board addressed the remaining asserted utilities, highlighting in particular the use of the claimed ESTs to monitor gene expression by measuring the level of mRNA through microarray technology and to serve as molecular markers. The Board found that using the claimed ESTs in screens does not provide a specific benefit because the application fails to provide any teaching regarding how to use the data relating to gene expression. *Id.*, slip op. at 21. The Board analogized the facts to those in *Brenner v. Manson*, 383 U.S. 519, 86 S.Ct. 1033, 16 L.Ed.2d 69 (1966), in which an applicant claimed a process of making a compound having no known use. In that case, the Supreme Court affirmed the rejection of the application on § 101 grounds. Here, the Board reasoned: “Just as the process in *Brenner* lacked utility because the specification did not disclose how to use the end-product, the products claimed here lack utility, because even if used in gene expression assays, the specification does not disclose how to use SEQ ID NO: 1–5 specific gene expression data.” *Id.*, slip op. at 22. The Board offered a similar rationale for the use of the claimed ESTs as molecular markers. *Id.*, slip op. at 24. Accordingly, the Board affirmed the examiner’s rejection of the ’643 application for lack of utility under § 101. The Board also affirmed the examiner’s rejection of the ’643 application for lack of enablement under § 112, first paragraph, since the enablement rejection was made as a corollary to the utility rejection.

Fisher timely appealed. We have jurisdiction over this appeal pursuant to 28 U.S.C. § 1295(a)(4) and 35 U.S.C. §§ 141 and 144.

II. DISCUSSION

[1] Whether an application discloses a utility for a claimed invention is a question of fact. *In re Ziegler*, 992 F.2d 1197, 1200 (Fed.Cir.1993). We consequently review the Board’s determination that the ’643 application failed to satisfy the utility requirement of § 101 for substantial evidence. *In re Gartside*, 203 F.3d 1305, 1315 (Fed.Cir.2000) (“Because our review of the Board’s decision is confined to the factual record compiled by the Board, we accordingly conclude that the ‘substantial evidence’ standard is appropriate for our review of Board factfindings.”).

A. Utility

1.

[2] Fisher asserts that the Board unilaterally applied a heightened standard for utility in the case of ESTs, conditioning patentability upon “some undefined ‘spectrum’ of knowledge concerning the corresponding gene function.” Fisher contends that the standard is not so high and that Congress intended the language of § 101 to be given broad construction. In particular, Fisher contends that § 101 requires only that the claimed invention “not be frivolous, or injurious to the well-being, good policy, or good morals of society,” essentially adopting Justice Story’s view of a useful invention from *Lowell v. Lewis*, 15 F.Cas. 1018, 1019 (No. 8568) (C.C.D.Mass. 1817). Under the correct application of the law, Fisher argues, the record shows that the claimed ESTs provide seven specific and substantial uses, regardless whether the functions of the genes corresponding to the claimed ESTs are known. Fisher claims that the Board’s attempt to

equate the claimed ESTs with the chemical compositions in *Brenner* was misplaced and that several decisions in the field of pharmaceuticals, namely, *Cross v. Iizuka*, 753 F.2d 1040 (Fed.Cir.1985), *Nelson v. Bowler*, 626 F.2d 853 (C.C.P.A.1980), and *In re Jolles*, 628 F.2d 1322 (C.C.P.A.1980), are analogous and support finding utility of the claimed ESTs. Fisher likewise argues that the general commercial success of ESTs in the marketplace confirms the utility of the claimed ESTs. Hence, Fisher avers that the Board's decision was not supported by substantial evidence and should be reversed.

The government agrees with Fisher that the utility threshold is not high, but disagrees with Fisher's allegation that the Board applied a heightened utility standard. The government contends that a patent applicant need disclose only a single specific and substantial utility pursuant to *Brenner*, the very standard articulated in the PTO's "Utility Examination Guidelines" ("Utility Guidelines") and followed here when examining the '643 application. It argues that Fisher failed to meet that standard because Fisher's alleged uses are so general as to be meaningless. What is more, the government asserts that the same generic uses could apply not only to the five claimed ESTs but also to any EST derived from any organism. It thus argues that the seven utilities alleged by Fisher are merely starting points for further research, not the end point of any research effort. It further disputes the importance of the commercial success of ESTs in the marketplace, pointing out that Fisher's evidence involved only databases, clone sets, and microarrays, not the five claimed ESTs. Therefore, the government

contends that we should affirm the Board's decision.

Several academic institutions and biotechnology and pharmaceutical companies³ write as amici curiae in support of the government. Like the government, they assert that Fisher's claimed uses are nothing more than a "laundry list" of research plans, each general and speculative, none providing a specific and substantial benefit in currently available form. The amici also advocate that the claimed ESTs are the objects of further research aimed at identifying what genes of unknown function are expressed during anthesis and what proteins of unknown function are encoded for by those genes. Until the corresponding genes and proteins have a known function, the amici argue, the claimed ESTs lack utility under § 101 and are not patentable.

[3] We agree with both the government and the amici that none of Fisher's seven asserted uses meets the utility requirement of § 101. Section 101 provides: "Whoever invents . . . any new and *useful* . . . composition of matter . . . may obtain a patent therefor . . ." (Emphasis added). In *Brenner*, the Supreme Court explained what is required to establish the usefulness of a new invention, noting at the outset that "a simple, everyday word ["useful," as found in § 101] can be pregnant with ambiguity when applied to the facts of life." 383 U.S. at 529, 86 S.Ct. 1033. Contrary to Fisher's argument that § 101 only requires an invention that is not "frivolous, injurious to the well-being, good policy, or good morals of society," the Supreme Court appeared to reject Justice Story's de minimis view of utility. *Id.* at 532-33, 86 S.Ct. 1033 (citation omitted).

3. Amici in support of the government include Affymetrix, Inc., American College of Medical Genetics, Association of American Medical Colleges, Baxter Healthcare Corporation,

Dow AgroSciences LLC, Eli Lilly and Company, Genentech, Inc., National Academy of Sciences, and the University of North Carolina School of Law.

The Supreme Court observed that Justice Story's definition "sheds little light on our subject," on the one hand framing the relevant inquiry as "whether the invention in question is 'frivolous and insignificant'" if narrowly read, while on the other hand "allowing the patenting of any invention not positively harmful to society" if more broadly read. *Id.* at 533, 86 S.Ct. 1033. In its place, the Supreme Court announced a more rigorous test, stating:

The basic *quid pro quo* contemplated by the Constitution and the Congress for granting a patent monopoly is the benefit derived by the public from an invention with *substantial utility*. Unless and until a process is refined and developed to this point—where *specific benefit exists in currently available form*—there is insufficient justification for permitting an applicant to engross what may prove to be a broad field.

Brenner, 383 U.S. at 534–35, 86 S.Ct. 1033 (emphases added). Following *Brenner*, our predecessor court, the Court of Customs and Patent Appeals, and this court have required a claimed invention to have a specific and substantial utility to satisfy § 101. See, e.g., *Fujikawa v. Wattanasin*, 93 F.3d 1559, 1563 (Fed.Cir.1996) ("Consequently, it is well established that a patent may not be granted to an invention unless substantial or practical utility for the invention has been discovered and disclosed.").

The Supreme Court has not defined what the terms "specific" and "substantial" mean per se. Nevertheless, together with the Court of Customs and Patent Appeals, we have offered guidance as to the uses which would meet the utility standard of § 101. From this, we can discern the kind of disclosure an application must contain to establish a specific and substantial utility for the claimed invention.

Courts have used the labels "practical utility" and "real world" utility interchangeably in determining whether an invention offers a "substantial" utility. Indeed, the Court of Customs and Patent Appeals stated that "[p]ractical utility" is a shorthand way of attributing 'real-world' value to claimed subject matter. In other words, one skilled in the art can use a claimed discovery in a manner which provides some *immediate benefit to the public*." *Nelson*, 626 F.2d at 856 (emphasis added).⁴ It thus is clear that an application must show that an invention is useful to the public as disclosed in its current form, not that it may prove useful at some future date after further research. Simply put, to satisfy the "substantial" utility requirement, an asserted use must show that that claimed invention has a significant and presently available benefit to the public.

Turning to the "specific" utility requirement, an application must disclose a use which is not so vague as to be meaningless. Indeed, one of our predecessor courts has observed "that the nebulous expressions 'biological activity' or 'biological properties' appearing in the specification convey no more explicit indication of the usefulness of the compounds and how to use them than did the equally obscure expression 'useful for technical and pharmaceutical purposes' unsuccessfully relied upon by the appellant in *In re Diedrich* [318 F.2d 946, 50 C.C.P.A. 1355 (1963)]." *In re Kirk*, 54 C.C.P.A. 1119, 376 F.2d 936, 941 (1967). Thus, in addition to providing a "substantial" utility, an asserted use must also show that that claimed invention can be used to provide a well-defined and particular benefit to the public.

4. In *Cross*, this court considered the phrase "practical utility" to be synonymous with the

phrase "substantial utility." 753 F.2d at 1047, n. 13.

[4] In 2001, partially in response to questions about the patentability of ESTs, the PTO issued Utility Guidelines governing its internal practice for determining whether a claimed invention satisfies § 101. See Utility Examination Guidelines, 66 Fed.Reg. 1092 (Jan. 5, 2001). The PTO incorporated these guidelines into the Manual of Patent Examining Procedure ("MPEP"). See U.S. Pat. & Trademark Off., Manual of Patent Examining Procedure § 2107 (8th ed.2001, rev. May 2004). The MPEP and Guidelines "are not binding on this court, but may be given judicial notice to the extent they do not conflict with the statute." *Enzo Biochem v. Gen-Probe*, 323 F.3d 956, 964 (Fed.Cir. 2002) (citing *Molins PLC v. Textron, Inc.*, 48 F.3d 1172, 1180 n. 10 (Fed.Cir.1995)). According to the Utility Guidelines, a specific utility is particular to the subject matter claimed and would not be applicable to a broad class of invention. Manual of Patent Examining Procedure § 2107.01. The Utility Guidelines also explain that a substantial utility defines a "real world" use. In particular, "[u]tilities that require or constitute carrying out further research to identify or reasonably confirm a 'real world' context of use are not substantial utilities." *Id.* Further, the Utility Guidelines discuss "research tools," a term often given to inventions used to conduct research. The PTO particularly cautions that

[a]n assessment that focuses on whether an invention is useful only in a research setting thus does not address whether the invention is in fact "useful" in a patent sense. [The PTO] must distinguish between inventions that have a specifically identified substantial utility and inventions whose asserted utility requires further research to identify or reasonably confirm.

Id. The PTO's standards for assessing whether a claimed invention has a specific and substantial utility comport with this

court's interpretation of the utility requirement of § 101.

Turning to the parties' arguments, Fisher first raises a legal issue, charging that the Board applied a heightened standard for utility in the case of ESTs. Fisher apparently bases this argument on statements made by the Board in connection with its discussion of whether the claimed ESTs can be used to identify a polymorphism. In that context, the Board stated:

Somewhere between having no knowledge (the present circumstances) and having complete knowledge of the gene and its role in the plant's development lies the line between 'utility' and 'substantial utility.' We need not draw the line or further define it in this case because the facts in this case represent the lowest end of the *spectrum*, i.e., an insubstantial use.

Board Decision, slip op. at 15 (emphasis added). Fisher reads the word "spectrum" out of context, claiming that the word somehow implies the application of a higher standard for utility than required by § 101. We conclude, however, that the Board did not apply an incorrect legal standard. In its decision, the Board made reference to a "spectrum" to differentiate between a substantial utility, which satisfies the utility requirement of § 101, and an insubstantial utility, which fails to satisfy § 101. The Board plainly did not announce or apply a new test for assessing the utility of ESTs. It simply followed the Utility Guidelines and MPEP, which mandate the specific and substantial utility test set forth in *Brenner*. Indeed, we note that Example 9 of the PTO's "Revised Interim Utility Guidelines Training Materials" is applicable to the facts here. See U.S. Pat. & Trademark Off., Revised Interim Utility Guidelines Training Materials 50-53 (1999), available at www.uspto.gov/web/menu/utility.pdf. In that example, a cDNA

fragment disclosed as being useful as a probe to obtain the full length gene corresponding to a cDNA fragment was deemed to lack a specific and substantial utility. Additionally, the MPEP particularly explains that a claim directed to a polynucleotide disclosed to be useful as a "gene probe" or "chromosome marker," as is the case here, fails to satisfy the specific utility requirement unless a specific DNA target is also disclosed. Manual of Patent Examining Procedure § 2107.01.

Regarding the seven uses asserted by Fisher, we observe that each claimed EST uniquely corresponds to the single gene from which it was transcribed ("underlying gene"). As of the filing date of the '643 application, Fisher admits that the underlying genes have no known functions. Fisher, nevertheless, claims that this fact is irrelevant because the seven asserted uses are not related to the functions of the underlying genes. We are not convinced by this contention. Essentially, the claimed ESTs act as no more than research intermediates that may help scientists to isolate the particular underlying protein-encoding genes and conduct further experimentation on those genes. The overall goal of such experimentation is presumably to understand the maize genome—the functions of the underlying genes, the identity of the encoded proteins, the role those proteins play during anthesis, whether polymorphisms exist, the identity of promoters that trigger protein expression, whether protein expression may be controlled, etc. Accordingly, the claimed ESTs are, in words of the Supreme Court, mere "object[s] of use-testing," to wit, objects upon which scientific research could be performed with no assurance that anything useful will be discovered in the end. *Brenner*, 383 U.S. at 535, 86 S.Ct. 1033.

Fisher compares the claimed ESTs to certain other patentable research tools, such as a microscope. Although this com-

parison may, on first blush, be appealing in that both a microscope and one of the claimed ESTs can be used to generate scientific data about a sample having unknown properties, Fisher's analogy is flawed. As the government points out, a microscope has the specific benefit of optically magnifying an object to immediately reveal its structure. One of the claimed ESTs, by contrast, can only be used to detect the presence of genetic material having the same structure as the EST itself. It is unable to provide any information about the overall structure let alone the function of the underlying gene. Accordingly, while a microscope can offer an immediate, real world benefit in a variety of applications, the same cannot be said for the claimed ESTs. Fisher's proposed analogy is thus inapt. Hence, we conclude that Fisher's asserted uses are insufficient to meet the standard for a "substantial" utility under § 101.

Moreover, all of Fisher's asserted uses represent merely hypothetical possibilities, objectives which the claimed ESTs, or any EST for that matter, *could* possibly achieve, but none for which they have been used in the real world. Focusing on the two uses emphasized by Fisher at oral argument, Fisher maintains that the claimed ESTs could be used to identify polymorphisms or to isolate promoters. Nevertheless, in the face of a utility rejection, Fisher has not presented any evidence, as the Board well noted, showing that the claimed ESTs have been used in either way. That is, Fisher does not present either a single polymorphism or a single promoter, assuming at least one of each exists, actually identified by using the claimed ESTs. Further, Fisher has not shown that a polymorphism or promoter so identified would have a "specific and substantial" use. The Board, in fact, correctly recognized this very deficiency and cited it

as one of the reasons for upholding the examiner's final rejection.

With respect to the remaining asserted uses, there is no disclosure in the specification showing that any of the claimed ESTs were used as a molecular marker on a map of the maize genome. There also is no disclosure establishing that any of the claimed ESTs were used or, for that matter, could be used to control or provide information about gene expression. Significantly, despite the fact that maize leaves produce over two thousand different proteins during anthesis, Fisher failed to show that one of the claimed ESTs translates into a portion of one of those proteins. Fisher likewise did not provide any evidence showing that the claimed ESTs were used to locate genetic molecules in other plants and organisms. What is more, Fisher has not proffered any evidence showing that any such generic molecules would themselves have a specific and substantial utility. Consequently, because Fisher failed to prove that its claimed ESTs can be successfully used in the seven ways disclosed in the '643 application, we have no choice but to conclude that the claimed ESTs do not have a "substantial" utility under § 101.

Furthermore, Fisher's seven asserted uses are plainly not "specific." Any EST transcribed from any gene in the maize genome has the potential to perform any one of the alleged uses. That is, any EST transcribed from any gene in the maize genome may be a molecular marker or a source for primers. Likewise, any EST transcribed from any gene in the maize genome may be used to measure the level of mRNA in a tissue sample, identify the presence or absence of a polymorphism, isolate promoters, control protein expression, or locate genetic molecules of other plants and organisms. Nothing about Fisher's seven alleged uses set the five claimed ESTs apart from the more than

32,000 ESTs disclosed in the '643 application or indeed from any EST derived from any organism. Accordingly, we conclude that Fisher has only disclosed general uses for its claimed ESTs, not specific ones that satisfy § 101.

We agree with the Board that the facts here are similar to those in *Brenner*. There, as noted above, the applicant claimed a process for preparing compounds of unknown use. Similarly, Fisher filed an application claiming five particular ESTs which are capable of hybridizing with underlying genes of unknown function found in the maize genome. The *Brenner* court held that the claimed process lacked a utility because it could be used only to produce a compound of unknown use. The *Brenner* court stated: "We find absolutely no warrant for the proposition that although Congress intended that no patent be granted on a chemical compound whose sole 'utility' consists of its potential role as an object of use-testing, a different set of rules was meant to apply to the process which yielded the unpatentable product." 383 U.S. at 535, 86 S.Ct. 1033. Applying that same logic here, we conclude that the claimed ESTs, which do not correlate to an underlying gene of known function, fail to meet the standard for utility intended by Congress.

In addition to approving of the Board's reliance on *Brenner*, we observe that the facts here are even more analogous to those presented in *Kirk*, 54 C.C.P.A. 1119, 376 F.2d 936, and *In re Joly*, 54 C.C.P.A. 1159, 376 F.2d 906 (1967), two cases decided by our predecessor court shortly after *Brenner*. In *Kirk*, the applicant sought to patent new steroidal compounds disclosed as having two possible utilities. First, the applicant alleged that the claimed compounds were useful for their "biological activity" because "one skilled in the art would know how to use the compounds ...

to take advantage of their presently-existing biological activity.” *Kirk*, 376 F.2d at 939. The court rejected this claimed utility on the ground that it was not sufficiently “specific,” but was instead “nebulous.” *Id.* at 941.

Second, the applicant asserted that the claimed compounds could be used by skilled chemists as intermediates in the preparation of final steroidal compounds of unknown use. Relying on *Brenner*, the court reasoned:

It seems clear that, if a process for producing a product of only conjectural use is not itself “useful” within § 101, it cannot be said that the starting materials for such a process—i.e., the presently claimed intermediates—are “useful.” It is not enough that the specification disclose that the intermediate exists and that it “works,” reacts, or can be used to produce some intended product of no known use. Nor is it enough that the product disclosed to be obtained from the intermediate belongs to some class of compounds which now is, or in the future might be, the subject of research to determine some *specific use*. Cf. *Reiners v. Mehlretter*, 43 C.C.P.A. 1019, 236 F.2d 418, 421 [(C.C.P.A.1956)] where compounds employed as intermediates to produce other directly useful compounds were found to be themselves useful.

Id. at 945–46 (emphasis added). Therefore, the court affirmed the Board’s rejection of the claimed compounds for lack of utility.

The facts in *Joly* are nearly identical to the facts in *Kirk*. The *Joly* applicant filed an application claiming compounds useful as intermediates in preparing steroids that were themselves not shown or known to be useful, but that were similar in chemical structure to steroids of known pharmacological usefulness. The court adopted the reasoning of the *Kirk* court in its entirety

and affirmed the Board’s decision rejecting the claimed intermediates for failing to comply with § 101. *Joly*, 376 F.2d at 908–09.

Just as the claimed compounds in *Kirk* and *Joly* were useful only as intermediates in the synthesis of other compounds of unknown use, the claimed ESTs can only be used as research intermediates in the identification of underlying protein-encoding genes of unknown function. The rationale of *Kirk* and *Joly* thus applies here. In the words of the *Kirk* court:

We do not believe that it was the intention of the statutes to require the Patent Office, the courts, or the public to play the sort of guessing game that might be involved if an applicant could satisfy the requirements of the statutes by indicating the usefulness of a claimed compound *in terms of possible use so general as to be meaningless* and then, after his research or that of his competitors has definitely ascertained an actual use for the compound, adducing evidence intended to show that a particular specific use would have been obvious to men skilled in the particular art to which this use relates.

376 F.2d at 942 (emphasis added).

That the *Kirk* and *Joly* decisions involved chemical compounds, while the present case involves biological entities, does not distinguish these decisions. The rationale presented therein, having been drawn from principles set forth by the Supreme Court in *Brenner*, applies with equal force in the fields of chemistry and biology as well as in any scientific discipline. In *Brenner*, the Supreme Court was primarily concerned with creating an unwarranted monopoly to the detriment of the public:

Whatever weight is attached to the value of encouraging disclosure and of inhibiting secrecy, we believe a more compel-

ling consideration is that a process patent in the chemical field, which has not been developed and pointed to the degree of specific utility, creates a monopoly of knowledge which should be granted only if clearly commanded by the statute. Until the process claim has been reduced to production of a product shown to be useful, the metes and bounds of that monopoly are not capable of precise delineation. It may engross a vast, unknown, and perhaps unknowable area. Such a patent may confer power to block off whole areas of scientific development, without compensating benefit to the public. . . . This is not to say that we mean to disparage the importance of contributions to the fund of scientific information short of the invention of something "useful," or that we are blind to the prospect that what now seems without "use" may tomorrow command the grateful attention of the public. But a patent is not a hunting license. It is not a reward for the search, but compensation for its successful conclusion. [A] patent system must be related to the world of commerce rather than to the realm of philosophy.

Brenner, 383 U.S. at 535-36, 86 S.Ct. 1033 (citations, quotation, and footnote omitted). Here, granting a patent to Fisher for its five claimed ESTs would amount to a hunting license because the claimed ESTs can be used only to gain further information about the underlying genes and the proteins encoded for by those genes. The claimed ESTs themselves are not an end of Fisher's research effort, but only tools to be used along the way in the search for a practical utility. Thus, while Fisher's claimed ESTs may add a noteworthy contribution to biotechnology research, our precedent dictates that the '643 application does not meet the utility requirement of § 101 because Fisher does not identify the function for the underlying protein-encoding genes. Absent such identification, we

hold that the claimed ESTs have not been researched and understood to the point of providing an immediate, well-defined, real world benefit to the public meriting the grant of a patent.

2.

Fisher's reliance on *Jolles*, *Nelson*, and *Cross*, cases which found utility in certain claimed pharmaceutical compounds, is misplaced. In *Jolles*, the applicant filed an application claiming naphthacene compounds useful in treating acute myeloblastic leukemia. To support the asserted utility, the applicant presented *in vivo* data showing eight of the claimed compounds effectively treated tumors in a mouse model. Our predecessor court reversed the Board's affirmance of the final rejection for lack of utility, finding that the structural similarity between the compounds tested *in vivo* and the remaining claimed compounds was sufficient to establish utility for the remaining claimed compounds. *Jolles*, 628 F.2d at 1327-28.

In *Nelson*, decided by the Court of Customs and Patent Appeals in the same year as *Jolles*, Nelson claimed prostaglandin compounds. The PTO declared an interference with an application filed by Bowler claiming the same compounds. The issue before the Board was whether Nelson had established utility for the claimed prostaglandins as smooth muscle stimulants and blood pressure modulators via *in vivo* and *in vitro* data, specifically, an *in vivo* rat blood pressure test and an *in vitro* gerbil colon smooth muscle stimulation test. The Board declined to award priority to Nelson, characterizing Nelson's tests as "rough screens, uncorrelated with actual utility [in humans]." Our predecessor court reversed, concluding that "tests evidencing pharmacological activity may manifest a practical utility even though they may not establish a specific therapeutic use." *Nelson*, 626 F.2d at 856.

In *Cross*, decided by the Federal Circuit five years after *Jolles* and *Nelson*, Iizuka filed an application claiming thromboxane synthetase inhibitors, alleged to be useful in treating inflammation, asthma, hypertension, and other ailments. When *Cross* filed an application claiming the same compounds two months after Iizuka, the PTO declared an interference. The dispositive issue concerned whether Iizuka's Japanese priority application disclosed utility for the claimed inhibitors. The Board concluded that it offered a sufficient disclosure based upon *in vitro* data showing strong inhibitory action for thromboxane synthetase for structurally-similar compounds in human or bovine platelet microsomes. We affirmed, reasoning:

Opinions of our predecessor court have recognized the fact that pharmacological testing of animals is a screening procedure for testing new drugs for practical utility. This *in vivo* testing is but an intermediate link in a screening chain which may eventually lead to the use of the drug as a therapeutic agent in humans. We perceive no insurmountable difficulty, under appropriate circumstances, in finding that the first link in the screening chain, *in vitro* testing, may establish a practical utility for the compound in question. Successful *in vitro* testing will marshal resources and direct the expenditure of effort to further *in vivo* testing of the most potent compounds, thereby providing an immediate benefit to the public, analogous to the benefit provided by the showing of an *in vivo* utility.

Cross, 753 F.2d at 1050 (citations omitted).

The facts in these three cases are readily distinguishable from the facts here. In *Jolles*, *Nelson*, and *Cross*, the applicants disclosed specific pharmaceutical uses in humans for the claimed compounds and supported those uses with specific animal test data, *in vitro*, *in vivo*, or both. In contrast, Fisher disclosed a variety of as-

serted uses for the claimed ESTs, but failed to present any evidence—test data, declaration, deposition testimony, or otherwise—to support those uses as presently beneficial and hence practical. Fisher did not show that even one of the claimed ESTs had been tested and successfully aided in identifying a polymorphism in the maize genome or in isolating a single promoter that could give clues about protein expression. Adopting the language of the *Cross* court, the alleged uses in *Jolles*, *Nelson*, and *Cross* were not “nebulous expressions, such as ‘biological activity’ or ‘biological properties’ [alleged in the application in *Kirk*],” that “convey little explicit indication regarding the utility of a compound.” *Cross*, 753 F.2d at 1048. Instead, the alleged uses in those cases gave a firm indication of the precise uses to which the claimed compounds could be put. For example, in *Nelson*, the claimed prostaglandins could be used to stimulate smooth muscle or modulate blood pressure in humans as shown by both *in vivo* and *in vitro* animal data. Hence, the *Jolles*, *Nelson*, and *Cross* courts concluded that the claimed pharmaceutical compounds satisfied the specific and substantial utility requirements of § 101. We cannot reach that same conclusion here. Fisher's laundry list of uses, like the terms “biological activity” or “biological properties” alleged in *Kirk*, are nebulous, especially in the absence of any data demonstrating that the claimed ESTs were actually put to the alleged uses.

Fisher's reliance on the commercial success of general EST databases is also misplaced because such general reliance does not relate to the ESTs at issue in this case. Fisher did not present any evidence showing that agricultural companies have purchased or even expressed any interest in the claimed ESTs. And, it is entirely unclear from the record whether such business entities ever will. Accordingly, while

commercial success may support the utility of an invention, it does not do so in this case. See *Raytheon Co. v. Roper Corp.*, 724 F.2d 951, 959 (Fed.Cir.1983) (stating that proof of a utility may be supported when a claimed invention meets with commercial success).

3.

As a final matter, we observe that the government and its amici express concern that allowing EST patents without proof of utility would discourage research, delay scientific discovery, and thwart progress in the "useful Arts" and "Science." See U.S. Const. art. I, § 8, cl. 8. The government and its amici point out that allowing EST claims like Fisher's would give rise to multiple patents, likely owned by several different companies, relating to the same underlying gene and expressed protein. Such a situation, the government and amici predict, would result in an unnecessarily convoluted licensing environment for those interested in researching that gene and/or protein.

The concerns of the government and amici, which may or may not be valid, are not ones that should be considered in deciding whether the application for the claimed ESTs meets the utility requirement of § 101. The same may be said for the resource and managerial problems that the PTO potentially would face if applicants present the PTO with an onslaught of patent applications directed to particular ESTs. Congress did not intend for these practical implications to affect the determination of whether an invention satisfies the requirements set forth in 35 U.S.C. §§ 101, 102, 103, and 112. They are public policy considerations which are more appropriately directed to Congress as the legislative branch of government, rather than this court as a judicial body responsible simply for interpreting and applying statutory law. Under Title 35, an applicant is entitled to a patent if his in-

vention is new, useful, nonobvious, and his application adequately describes the claimed invention, teaches others how to make and use the claimed invention, and discloses the best mode for practicing the claimed invention. What is more, when Congress enacted § 101, it indicated that "anything under the sun that is made by man" constitutes potential subject matter for a patent. S.Rep. No. 82-1979, at 7 (1952), U.S.Code Cong. & Admin.News at 2394, 2399. Policy reasons aside, because we conclude that the utility requirement of § 101 is not met, we hold that Fisher is not entitled to a patent for the five claimed ESTs.

B. Enablement

[5] Fisher asserts that we should reverse the enablement rejection upheld by the Board since the Board made it contingent upon the utility rejection, which Fisher argues was not supported by substantial evidence for reasons analyzed above. The government argues to the contrary, asserting that claim 1 of the '643 application cannot be enabled because the claimed ESTs were not disclosed as having a specific and substantial utility. We agree with the government. It is well established that the enablement requirement of § 112 incorporates the utility requirement of § 101.

The how to use prong of section 112 incorporates as a matter of law the requirement of 35 U.S.C. § 101 that the specification disclose as a matter of fact a practical utility for the invention. If the application fails as a matter of fact to satisfy 35 U.S.C. § 101, then the application also fails as a matter of law to enable one of ordinary skill in the art to use the invention under 35 U.S.C. § 112.

Ziegler, 992 F.2d at 1200-01 (citations omitted); see also *Kirk*, 376 F.2d at 942 ("Necessarily, compliance with § 112 re-

quires a description of how to use presently useful inventions, otherwise an applicant would anomalously be required to teach how to use a useless invention.”); *In re Brana*, 51 F.3d 1560, 1564 (Fed.Cir.1995) (“Obviously, if a claimed invention does not have utility, the specification cannot enable one to use it.”); Manual of Patent Examining Procedure § 2107.01. Here, in light of our conclusion that the Board’s decision with respect to utility applied the correct legal standard and was supported by substantial evidence, we conclude that Fisher failed to satisfy the enablement requirement. Consequently, we leave undisturbed the enablement rejection of the ’643 application under § 112, first paragraph.

III. CONCLUSION

We conclude that substantial evidence supports the Board’s findings that each of the five claimed ESTs lacks a specific and substantial utility and that they are not enabled. Accordingly, the Board’s decision affirming the final rejection of claim 1 of the ’643 patent for lack of utility under § 101 and lack of enablement under § 112, first paragraph, is affirmed.

AFFIRMED.

RADER, Circuit Judge, dissenting.

This court today determines that expressed sequence tags (ESTs) do not satisfy 35 U.S.C. § 101 unless there is a known use for the genes from which each EST is transcribed. While I agree that an invention must demonstrate utility to satisfy § 101, these claimed ESTs have such a utility, at least as research tools in isolating and studying other molecules. Therefore, I respectfully dissent.

Several, if not all, of Fisher’s asserted utilities claim that ESTs function to study other molecules. In simple terms, ESTs are research tools. Admittedly ESTs have use only in a research setting. However, the value and utility of research tools gen-

erally is beyond question, even though limited to a laboratory setting. See U.S. Pat. & Trademark Off., Manual of Patent Examining Procedure (MPEP) § 2107.01 at 2100–33 (8th ed.2001, rev.Feb.2003) (“Many research tools such as gas chromatographs, screening assays, and nucleotide sequencing techniques have a clear, specific and unquestionable utility (e.g., they are useful in analyzing compounds).”). Thus, if the claimed ESTs qualify as research tools, then they have a “specific” and “substantial” utility sufficient for § 101. If these ESTs do not enhance research, then *Brenner v. Manson*, 383 U.S. 519, 86 S.Ct. 1033, 16 L.Ed.2d 69 (1966) (involving the patentability of methods for producing compounds having no known use) controls and erects a § 101 bar for lack of utility. For the following reasons, these claimed ESTs are more akin to patentable research tools than to the unpatentable methods in *Brenner*.

In *Brenner*, the Court confronted a growing conflict between this court’s predecessor, the Court of Customs and Patent Appeals (CCPA), and the Patent Office over the patentability of methods of producing compounds with no known use. This conflict began with *In re Nelson*, 47 C.C.P.A. 1031, 280 F.2d 172 (1960), the first in a series of cases wherein the CCPA reversed several Patent Office utility rejections. *Brenner*, 383 U.S. at 530, 86 S.Ct. 1033. *Brenner* put an end to these cases because, in the 1960s, the Court could not distinguish between denying patents to compounds with no known use and denying patents to methods of producing those useless compounds. The Court commented:

We find absolutely no warrant for the proposition that although Congress intended that no patent be granted on a chemical compound whose sole ‘utility’ consists of its potential role as an object of use-testing, a different set of rules

was meant to apply to the process which yielded the unpatentable product. That proposition seems to us little more than an attempt to evade the impact of the rules which concededly govern patentability of the product itself.

Id. at 535, 86 S.Ct. 1033. This court's predecessor later extended *Brenner* to bar patents on compounds as intermediates in the preparation of other compounds having no known use. See *In re Kirk*, 54 C.C.P.A. 1119, 376 F.2d 936 (1967) (rejecting intermediaries for steroids with no known use). These cases, however, share a common underpinning—a method of producing a compound with no known use has no more benefit to society than the useless compound itself.

This case is very different. Unlike the methods and compounds in *Brenner* and *Kirk*, Fisher's claimed EST's are beneficial to society. As an example, these research tools "may help scientists to isolate the particular underlying protein-encoding genes... [with the] overall goal of such experimentation... presumably [being] to understand the maize genome[.]" *Majority Opinion*, at 1373. They also can serve as a probe introduced into a sample tissue to confirm "that the gene corresponding to the EST was being expressed in the sample tissue at the time of mRNA extraction." *Id.*, at 1367.

These research tools are similar to a microscope; both take a researcher one step closer to identifying and understanding a previously unknown and invisible structure. Both supply information about a molecular structure. Both advance research and bring scientists closer to unlocking the secrets of the corn genome to provide better food production for the hungry world. If a microscope has \$ 101 utility, so too do these ESTs.

The Board and this court acknowledge that the ESTs perform a function, that they have a utility, but proceed quickly to

a value judgment that the utility would not produce enough valuable information. The Board instead complains that the information these ESTs supply is too "insubstantial" to merit protection. Yet this conclusion denies the very nature of scientific advance. Science always advances in small incremental steps. While acknowledging the patentability of research tools generally (and microscopes as one example thereof), this court concludes with little scientific foundation that these ESTs do not qualify as research tools because they do not "offer an immediate, real world benefit" because further research is required to understand the underlying gene. This court further faults the EST research for lacking any "assurance that anything useful will be discovered in the end." These criticisms would foreclose much scientific research and many vital research tools. Often scientists embark on research with no assurance of success and knowing that even success will demand "significant additional research."

Nonetheless, this court, oblivious to the challenges of complex research, discounts these ESTs because it concludes (without scientific evidence) that they do not supply enough information. This court reasons that a research tool has a "specific" and "substantial" utility *only* if the studied object is readily understandable using the claimed tool—that no further research is required. Surely this cannot be the law. Otherwise, only the final step of a lengthy incremental research inquiry gets protection.

Even with a microscope, significant additional research is often required to ascertain the particular function of a "revealed" structure. To illustrate, a cancerous growth, magnified with a patented microscope, can be identified and distinguished from other healthy cells by a properly trained doctor or researcher. But even today, the scientific community

still does not fully grasp the reasons that cancerous growths increase in mass and spread throughout the body,¹ or the nature of compounds that interact with them, or the interactions of environmental or genetic conditions that contribute to developing cancer. Significant additional research is required to answer these questions. Even with answers to these questions, the cure for cancer will remain in the distance. Yet the microscope still has “utility” under § 101. Why? Because it takes the researcher one step closer to answering these questions. Each step, even if small in isolation, is nonetheless a benefit to society sufficient to give a viable research tool “utility” under § 101. In fact, experiments that fail still serve to eliminate some possibilities and provide information to the research process.

The United States Patent Office, above all, should recognize the incremental nature of scientific endeavor. Yet, in the interest of easing its administrative load, the Patent Office will eliminate some research tools as providing “insubstantial” advances. How does the Patent Office know which “insubstantial” research step will contribute to a substantial breakthrough in genomic study? Quite simply, it does not.

In addition, this court faults Fisher for not presenting evidence of utility showing that the claimed ESTs “have been used in the real world.” To the contrary, this court misapprehended the proper procedure. Fisher asserted seven different utilities. The Board rejected two of these assertions outright as “insubstantial.” See *Ex parte Fisher*, App. No.2002-2046, slip. op at 14-16 (Bd. Pat.App. and Int.2004) (acknowledging that the ESTs may be able

to detect “the absence of a polymorphism” and “to isolate nucleic acid molecules of other plants and organisms[,]” but finding such utilities are not “substantial” even if the ESTs can perform them). This summary dismissal deprived Fisher of any chance to proffer evidence. Rather than fault Fisher for not presenting evidence it was prevented from offering, this court should instead observe that the Board did not satisfy its burden of challenging Fisher’s presumptively correct assertion that the ESTs were *capable* of performing those functions. See MPEP § 2107.02(IV) at 2100-40 (noting that the initial burden is on the office to establish a prima facie case as to lack of utility and to provide evidentiary support thereof); *In re Brana*, 51 F.3d 1560, 1566 (Fed.Cir.1995) (where an applicant has asserted utility in the disclosure, the Patent Office has the initial burden of challenging this presumptively correct assertion of utility).

Abandoning the proper legal procedure, the Board reasoned that the molecules studied with these ESTs showed no particular use, therefore the ESTs themselves also lacked a utility. In so ruling, the Board did not reject Fisher’s utilities on the basis that the ESTs were *unable* to perform the purported utilities. Thus, the Board did not establish a prima facie challenge to the ESTs’ ability to perform these two utilities. Without anything to rebut, Fisher had no obligation or opportunity to provide evidence in rebuttal. Thus, I respectfully disagree with this court’s conclusion that the Board’s decision can be affirmed on the basis that Fisher did not supply evidence of the ESTs’ ability to perform the asserted utilities.

In truth, I have some sympathy with the Patent Office’s dilemma. The Office needs

1. ESTs have already been used to advance cancer research well beyond what is achievable using microscopes alone. See Andy J. Minn, *Genes That Mediate Breast Cancer Me-*

tastasis To Lung, Nature, July 28, 2005 at 518-24 (discussing research to identify genes that mark and mediate breast cancer metastasis to the lung).

some tool to reject inventions that may advance the "useful arts" but not sufficiently to warrant the valuable exclusive right of a patent. The Patent Office has seized upon this utility requirement to reject these research tools as contributing "insubstantially" to the advance of the useful arts. The utility requirement is ill suited to that task, however, because it lacks any standard for assessing the state of the prior art and the contributions of the claimed advance. The proper tool for assessing sufficient contribution to the useful arts is the obviousness requirement of 35 U.S.C. § 103. Unfortunately this court has deprived the Patent Office of the obviousness requirement for genomic inventions. See *In re Deuel*, 51 F.3d 1552 (Fed. Cir.1995); Martin J. Adelman et al., *Patent Law*, 517 (West Group 1998) (commenting that scholars have been critical of *Deuel*, which "overly favored patent applicants in biotech by adopting an overly lax nonobviousness standard." (citing Anita Varma & David Abraham, *DNA Is Different: Legal Obviousness and the Balance Between Biotech Inventors and the Market*, 9 Harv. J.L. & Tech. 53 (1996))); Philippe Ducor, *The Federal Circuit and In re Deuel: Does § 103 apply to Naturally Occurring DNA?*, 77 J. Pat. & Trademark Off. Soc'y 871, 883 (Nov.1995) ("The Court of Appeals for the Federal Circuit could have formulated its opinion in only one

sentence: '35 U.S.C. § 103 does not apply to newly retrieved natural DNA sequences.'"); Philippe Ducor, *Recombinant Products and Nonobviousness: A Typology*, 13 Santa Clara Computer and High Tech. L.J. 1, 44-45 (Feb.1997) ("This amounts to a practical elimination of the requirement for nonobviousness for these products, even when all the information necessary to discover them is previously available."); see also over fifty additional articles critical of *Deuel* in the "Citing References" tab for *Deuel* on Westlaw. Nonetheless, rather than distort the utility test, the Patent Office should seek ways to apply the correct test, the test used world wide for such assessments (other than in the United States), namely inventive step or obviousness.

Thus, for the foregoing reasons, I would find that Fisher's asserted utilities qualify the claimed ESTs as research tools useful in the study of other molecules. Because research tools provide a cognizable benefit to society, much like a microscope, the ESTs claimed here have "utility" under § 101. In addition, the enablement rejection should also be reversed because it was a consequence of the finding of lack of utility.

